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November 24, 1998

**Box Patent Application**

Assistant Commissioner for Patents  
Washington, D.C. 20231

Re: U.S. Non-Provisional Patent Application  
Appl. No.: To Be Assigned  
Filed: November 24, 1998  
For: **NUCLEIC ACID MOLECULES AND OTHER MOLECULES  
ASSOCIATED WITH THE METHIONINE SYNTHESIS AND  
DEGRADATION PATHWAYS**  
Inventor(s): Stefan A. Bledig *et al.*  
Our Ref: 04983.0002US01/38-21 (15077)B

Sir:

The following documents are being submitted under 37 C.F.R. § 1.53(b)(2) herewith for appropriate action by the U.S. Patent and Trademark Office:

1. Patent Application Utility Application;
2. Utility Patent Application Transmittal (PTO/SB/05);
3. Form PTO-1082 (in duplicate);
4. U.S. Utility Patent Application entitled

**Nucleic Acid Molecules And Other Molecules Associated With The  
Methionine Synthesis And Degradation Pathways**

and naming as inventor(s):

**Stefan A. Bledig, Joseph R. Byrum, Gregory J. Hinkle and JingDong  
Liu**

the application consisting of:

- (i) 288 pages of description prior to the claims;
  - (ii) 5 pages of claims (11 claims);
  - (iii) a one (1) page abstract;
  - (iv) sequence listing of 1127 pages;
- 5. Sequence Listing Disk;
  - 6. Statement Regarding Sequence Submissions;
  - 7. Information Disclosure Statement;
  - 8. Form PTO-1449 (9 pages) with 30 accompanying documents; and
  - 9. Two (2) return postcards.

It is respectfully requested that, of the two attached postcards, one be stamped with the filing date of these documents and returned to our courier, and the other, prepaid postcard, be stamped with the filing date and returned as soon as possible.

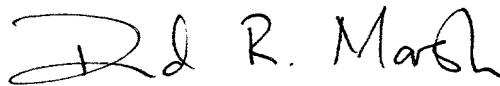
The present application claims priority under 35 U.S.C § 119(e) to provisional applications No. 60/067000 filed November 24, 1997, No. 60/066,873 filed November 25, 1997, No. 60/069472 filed December 09, 1997, No. 60/074,201 filed February 10, 1998, No. 60/074282 filed February 10, 1998, No. 60/074280 filed February 10, 1998, No. 60/074281 filed February 10, 1998, No. 60/074566 filed February 12, 1998, No. 60/074567 filed February 12, 1998, No. 60/074565 filed February 12, 1998, No. 60/075462 filed February 19, 1998, No. 60/074789 filed February 19, 1998, No. 60/075459 filed February 19, 1998, No. 60/075461 filed February 19, 1998, No. 60/075464 filed February 19, 1998, No. 60/075460 filed February 19, 1998, No. 60/075463 filed February 19, 1998, No. 60/077231 filed March 09, 1998, No. 60/077229 filed March 09, 1998, No. 60/077230 filed March 09, 1998, No. 60/078031 filed March 16, 1998, No. 60/078368 filed March 18, 1998, No. 60/080844 filed April 07, 1998, No. 60/083067 filed April 27, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15348)A filed April 29, 1998, No. 60/083387 filed April 29, 1998, No. 60/083388 filed April 29, 1998, No. 60/083389 filed April 29, 1998, "Nucleic Acid Molecules and Other Molecules Associated with the Ethylene Biosynthetic Pathway" docket No. 04983.0018/38-21(15097)A filed May 08, 1998, No. 60/085,245 filed May 13, 1998, No.



60/085224 filed May 13, 1998, No. 60/085223 filed May 13, 1998, No. 60/085222 filed May 13, 1998, No. 60/086186 filed May 21, 1998, No. 60/086,339 filed May 21, 1998, No. 60/086187 filed May 21, 1998, No. 60/086185 filed May 21, 1998, No. 60/086184 filed May 21, 1998, No. 60/086183 filed May 21, 1998, No. 60/086188 filed May 21, 1998, No. 60/089,524 filed June 16, 1998, No. 60/089,810 filed June 18, 1998, No. 60/089,814 filed June 18, 1998, "Nucleic acid molecules and other molecules associated with the Plant Sugar and Nitrogen Transporters Pathway" docket No. 04983.0043/38-21(15412)A filed June 30, 1998, No. 60/092,036 filed July 08, 1998, No. 60/099667 filed September 09, 1998, No. 60/099668 filed September 09, 1998, No. 60/099670 filed September 09, 1998, No. 60/099697 filed September 09, 1998, No. 60/100674 filed September 16, 1998, No. 60/100673 filed September 16, 1998, No. 60/100672 filed September 16, 1998, No. 60/101132 filed September 21, 1998, No. 60/101130 filed September 21, 1998, "Nucleic acid molecules and other molecules associated with Plants" docket No. 38-21(15459)A filed September 21, 1998, No. 60/101344 filed September 22, 1998, No. 60/101347 filed September 22, 1998, No. 60/101343 filed September 22, 1998, No. 60/104,126 filed October 13, 1998, No. 60/104,128 filed October 13, 1998, No. 60/104,127 filed October 13, 1998, No. 60/104,124 filed October 13, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15445)A filed November 18, 1998 and "Nucleic Acid Molecules and other Molecules associated with Plants" docket No. 38-21(15592) filed November 18, 1998 hereby incorporated by reference herein in their entirety.

This application is being filed under 37 C.F.R. § 1.53(b) without Declaration and without filing fee.

Respectfully submitted,



David R. Marsh (Reg. No. 41,408)

Kevin W. McCabe (Reg. No. 41,182)

Enclosures

Please type a plus sign (+) inside this box → ☐

<b>UTILITY PATENT APPLICATION TRANSMITTAL</b> <small>(Only for new nonprovisional applications under 37 CFR 1.53(h))</small>		Attorney Docket No. 04983.0002US01/38-21(15077)B	
		First Named Inventor or Application Identifier Stefan A. Bledig	
		Title Nucleic Acid Molecules and Other Molecules Associated with the Methionine Synthesis and Degradation Pathways	
		Express Mail Label No.	

<b>APPLICATION ELEMENTS</b> <i>See MPEP chapter 600 concerning utility patent application contents</i>	<b>ADDRESS TO:</b> Assistant Commissioner for Patents Box Patent Application Washington, DC 20231
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<p>1. <input checked="" type="checkbox"/> Fee Transmittal Form (Form PTO-1082) <i>(Submit an original and a duplicate for fee processing)</i></p> <p>2. <input checked="" type="checkbox"/> Specification [Total Pages 294 ] <i>(preferred arrangement set forth below)</i> - Descriptive title of the Invention - Cross References to Related Applications - Statement Regarding Fed sponsored R&amp;D - Reference to Microfiche Appendix - Background of the Invention - Brief Summary of the Invention - Brief Description of the Drawings (if filed) - Detailed Description - Claims - Abstract of the Disclosure</p> <p>3. <input type="checkbox"/> Drawing(s) (35 USC 113) [Total Sheets ]</p> <p>4. <input type="checkbox"/> Oath or Declaration [Total Pages ]  <input type="checkbox"/> Newly executed (original or copy)  <input type="checkbox"/> Copy from a prior application (37 CFR 1.63(d)) <i>(for continuation/divisional with Box 17 completed)</i> <i>[Note Box 5 below]</i>  i. <input type="checkbox"/> <b>DELETION OF INVENTOR(S)</b> Signed statement attached deleting inventor(s) named in the prior application, see 37 CFR 1.63(d)(2) and 1.33(b)</p> <p>5. <input type="checkbox"/> Incorporation By Reference <i>(useable if Box 4b is checked)</i> The entire disclosure of the prior application, from which a copy of the oath or declaration is supplied under Box 4b, is considered as being part of the disclosure of the accompanying application and is hereby incorporated by reference therein.</p>	<p>6. <input type="checkbox"/> Microfiche Computer Program <i>(Appendix)</i></p> <p>7. Nucleotide and/or Amino Acid Sequence Submission <i>(if applicable, all necessary)</i> a. <input checked="" type="checkbox"/> Computer Readable Copy b. <input checked="" type="checkbox"/> Paper Copy (identical to computer copy) c. <input checked="" type="checkbox"/> Statement verifying identity of above copies</p>
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<b>ACCOMPANYING APPLICATION PARTS</b>	
8. <input type="checkbox"/> Assignment Papers (cover sheet & document(s)) 9. <input type="checkbox"/> 37 CFR 3.73(b) Statement <input type="checkbox"/> Power of Attorney <i>(when there is an assignee)</i> 10. <input type="checkbox"/> English Translation Document <i>(if applicable)</i> 11. <input checked="" type="checkbox"/> Information Disclosure Statement (IDS)/PTO-1449 <input checked="" type="checkbox"/> Copies of IDS Citations 12. <input type="checkbox"/> Preliminary Amendment 13. <input checked="" type="checkbox"/> Return Receipt Postcard (MPEP 503) (Two) <i>(should be specifically itemized)</i> 14. <input type="checkbox"/> *Small Entity Statement(s) <input type="checkbox"/> Statement filed in prior application, Status still proper and desired 15. <input type="checkbox"/> Certified Copy of Priority Document(s) <i>(if foreign priority is claimed)</i> 16. <input type="checkbox"/> Other:	

\*NOTE FOR ITEMS 1 & 14 IN ORDER TO BE ENTITLED TO PAY SMALL ENTITY FEES, A SMALL ENTITY STATEMENT IS REQUIRED (37 C.F.R. § 1.27). EXCEPT IF ONE FILED IN A PRIOR APPLICATION IS RELIED UPON (37 C.F.R. § 1.28)

17. If a <b>CONTINUING APPLICATION</b> , check appropriate box and supply the requisite information: <input type="checkbox"/> Continuation <input type="checkbox"/> Divisional <input type="checkbox"/> Continuation-in-part (CIP) of prior application No: / Prior Application Information: Examiner: Group/Art Unit:	
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<b>18. CORRESPONDENCE ADDRESS</b>					
<input type="checkbox"/> Customer Number or Bar Code Label or <input checked="" type="checkbox"/> Correspondence address below <i>(Insert Customer No. or Attach bar code label here)</i>					
NAME		HOWREY & SIMON			
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Attorney Docket No. 04983.0002US01/38-21 (15077)B

ASSISTANT COMMISSIONER FOR PATENTS  
 Washington, D.C. 20231

Sir:

Transmitted herewith for filing is the patent application of  
 Inventors: Stefan A. Bledig *et al.*

For: **Nucleic Acid Molecules and Other Molecules Associated with the Methionine Synthesis and Degradation Pathways**

The filing fee has been calculated as shown below:

(Col. 1)		(Col. 2)	SMALL ENTITY		OR	OTHER THAN A SMALL ENTITY	
FOR	NO. FILED	NO EXTRA	RATE	FEE		RATE	FEE
BASIC FEE				\$ 380.00	OR		\$ 760.00
TOTAL CLAIMS	-20 =	*	x 9 =		OR	x 18 =	
INDEP. CLAIMS	-3 =	*	x 39 =		OR	x 78 =	
MULTIPLE DEPENDENT CLAIM PRESENTED			+ 130 =		OR	+ 260 =	
			TOTAL		OR	TOTAL	

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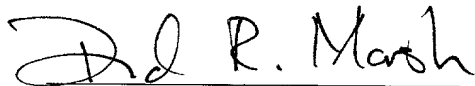
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- Any patent application processing fees under 37 CFR 1.17  
 — The issue fee set in 37 CFR 1.18 at or before mailing of the Notice of Allowance, pursuant to 37 CFR 1.311(b).  
 — Any filing fees under 37 CFR 1.16 for presentation of extra claims.

Date November 24, 1998



David R. Marsh (Reg. No. 41,408)  
 Kevin W. McCabe (Reg. No. 41,182)

# **NUCLEIC ACID MOLECULES AND OTHER MOLECULES ASSOCIATED WITH**

## **THE METHIONINE SYNTHESIS AND DEGRADATION PATHWAYS**

### **CROSS REFERENCE TO RELATED APPLICATIONS**

This application claims the benefit of applications No. 60/067000 filed November 24, 1997, No. 60/066,873 filed November 25, 1997, No. 60/069472 filed December 09, 1997, No. 60/074,201 filed February 10, 1998, No. 60/074282 filed February 10, 1998, No. 60/074280 filed February 10, 1998, No. 60/074281 filed February 10, 1998, No. 60/074566 filed February 12, 1998, No. 60/074567 filed February 12, 1998, No. 60/074565 filed February 12, 1998, No. 60/075462 filed February 19, 1998, No. 60/074789 filed February 19, 1998, No. 60/075459 filed February 19, 1998, No. 60/075461 filed February 19, 1998, No. 60/075464 filed February 19, 1998, No. 60/075460 filed February 19, 1998, No. 60/075463 filed February 19, 1998, No. 60/077231 filed March 09, 1998, No. 60/077229 filed March 09, 1998, No. 60/077230 filed March 09, 1998, No. 60/078031 filed March 16, 1998, No. 60/078368 filed March 18, 1998, No. 60/080844 filed April 07, 1998, No. 60/083067 filed April 27, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15348)A filed April 29, 1998, No. 60/083387 filed April 29, 1998, No. 60/083388 filed April 29, 1998, No. 60/083389 filed April 29, 1998, "Nucleic Acid Molecules and Other Molecules Associated with the Ethylene Biosynthetic Pathway" docket No. 04983.0018/38-21(15097)A filed May 08, 1998, No. 60/085,245 filed May 13, 1998, No. 60/085224 filed May 13, 1998, No. 60/085223 filed May 13, 1998, No. 60/085222 filed May 13, 1998, No. 60/086186 filed May 21, 1998, No. 60/086,339 filed May 21, 1998, No. 60/086187 filed May 21, 1998, No. 60/086185 filed May 21, 1998, No. 60/086184 filed May 21, 1998, No. 60/086183 filed May 21, 1998, No. 60/086188 filed May 21, 1998, No. 60/089,524 filed June 16, 1998, No. 60/089,810 filed June 18, 1998, No.

60/089,814 filed June 18, 1998, "Nucleic acid molecules and other molecules associated with the Plant Sugar and Nitrogen Transporters Pathway" docket No. 04983.0043/38-21(15412)A filed June 30, 1998, No. 60/092,036 filed July 08, 1998, No. 60/099667 filed September 09, 1998, No. 60/099668 filed September 09, 1998, No. 60/099670 filed September 09, 1998, No. 60/099697 filed September 09, 1998, No. 60/100674 filed September 16, 1998, No. 60/100673 filed September 16, 1998, No. 60/100672 filed September 16, 1998, No. 60/101132 filed September 21, 1998, No. 60/101130 filed September 21, 1998, "Nucleic acid molecules and other molecules associated with Plants" docket No. 38-21(15459)A filed September 21, 1998, No. 60/101344 filed September 22, 1998, No. 60/101347 filed September 22, 1998, No. 60/101343 filed September 22, 1998, No. 60/104,126 filed October 13, 1998, No. 60/104,128 filed October 13, 1998, No. 60/104,127 filed October 13, 1998, No. 60/104,124 filed October 13, 1998, "Nucleic Acid Molecules and Other Molecules Associated with Plants" docket No. 38-21(15445)A filed November 18, 1998 and "Nucleic Acid Molecules and other Molecules associated with Plants" docket No. 38-21(15592) filed November 18, 1998 hereby incorporated by reference herein in their entirety.

### **FIELD OF THE INVENTION**

The present invention is in the field of plant biochemistry. More specifically the invention relates to nucleic acid sequences from plant cells, in particular, DNA sequences from maize and soybean plants associated with the methionine pathway. The invention encompasses nucleic acid molecules that encode proteins and fragments of proteins. In addition, the invention also encompasses proteins and fragments of proteins so encoded and antibodies capable of binding these proteins or fragments. The invention also relates to methods of using the nucleic acid molecules, proteins and fragments of proteins and antibodies, for example for genome

mapping, gene identification and analysis, plant breeding, preparation of constructs for use in plant gene expression and transgenic plants.

## **BACKGROUND OF THE INVENTION**

### **I. METHIONINE SYNTHESIS PATHWAY**

The amino acid, L-methionine, is synthesized in higher plants via a pathway that starts with L-aspartate. This pathway has been studied (Azevedo *et al.*, *Phytochemistry* 46:395-419 (1997), the entirety of which is herein incorporated by reference). L-methionine is one of four so-called aspartate-derived amino acids (along with L-lysine, L-threonine and L-isoleucine)(Mifflin *et al.*, In: *Nitrogen Assimilation in Plants*, Hewitt *et al.*, (eds.), Academic Press, New York, 335 (1997); Bryan, In: *The Biochemistry of Plants*, Mifflin (ed.), Academic Press, New York, 403 (1980); Lea *et al.*, In: *The Chemistry and Biochemistry of Amino Acids*, Barrett *et al.*, (eds.), London, 5:197 (1985); Bryan, In: *The Biochemistry of Plants*, Mifflin *et al.*, (eds.), Academic Press, San Diego, 16:161 (1990), all of which are herein incorporated by reference in their entirety).

The methionine-specific part of the aspartate pathway includes the following enzymes: aspartate kinase (EC 2.7.2.4), aspartate-semialdehyde dehydrogenase (EC 1.2.1.11), homoserine dehydrogenase (EC 1.1.1.3), homoserine kinase (EC 2.7.1.39), cystathionine  $\gamma$ -synthase (EC 4.2.99.9), cystathionine  $\beta$ -lyase (EC 4.4.1.8) and methionine synthase (EC 2.1.1.14).

Aspartate kinase catalyzes the first reaction of the pathway in which aspartate is converted to  $\beta$ -aspartyl phosphate. This enzyme has been isolated and characterized from plant sources including maize, barley, carrot, pea and soybean. These studies have revealed that there are multiple isoenzymes of aspartate kinase and the isoenzymes differ with respect to both

feedback inhibition sensitivity and expression profile (tissue and developmental stage).

Feedback inhibition is mediated by lysine and threonine. Transgenic plants which express an unregulated aspartate kinase have demonstrated increased flux through the aspartate pathway.

Pathway regulation is reported to be exerted, at least in part, via control of this enzyme's activity.

Aspartate semialdehyde dehydrogenase catalyses the second pathway reaction and converts  $\beta$ -aspartyl phosphate to aspartate semialdehyde via an NADPH-dependent reaction.

Gengenbach *et al.*, *Crop Science* 18:472-476 (1978), the entirety of which is herein incorporated by reference, report the isolation of aspartate semialdehyde dehydrogenase from maize suspension culture cells. These suspension cultures did not exhibit feedback inhibition of the enzyme in the presence of aspartate-derived amino acids, with the exception of methionine, for which some feedback sensitivity was observed. Aspartate semialdehyde dehydrogenase enzyme activity has been reported in maize shoot, maize root and maize kernel (Gengenbach *et al.*, *Crop Science* 18:472-476 (1978)).

Homoserine dehydrogenase catalyzes the next step of the pathway in which homoserine is generated from aspartate semialdehyde in a reaction requiring NADH or NADPH.

Homoserine dehydrogenase enzyme has been studied in higher plants and multiple isoenzyme forms have been reported (Bryan *et al.*, *Biochemistry and Biophysics Research Communications* 41:1211-1217 (1970); Gengenbach *et al.*, *Crop Science* 18:472-476 (1978); Dotson *et al.*, *Plant Physiology* 91:1602-1608 (1989); Dotson *et al.*, *Plant Physiology* 93:98-104 (1989); Azevedo *et al.*, *Phytochemistry* 31:3725-3730 (1992); Azevedo *et al.*, *Phytochemistry* 31:3731-3734 (1992); Brennecke *et al.*, *Phytochemistry* 41:707 (1996); Aarnes, *Plant Science Letters* 9:137-145 (1977); Bright *et al.*, *Biochemical Genetics* 200:229-243 (1982); Aruda *et al.*, *Plant Physiology*

76:442-446 (1984); Lea *et al.*, In: *Barley: Genetics, Molecular Biology and Biotechnology* Shewrey (ed.), CAB International, Oxford 181 (1992); Davies *et al.*, *Plant Science Letters* 9:323-332 (1977); Davies *et al.*, *Plant Physiology* 62:536-541 (1978); Matthews *et al.*, *Zeitschrift für Naturforschung, Section Bioscience* 34:1177-1185 (1979); Relton *et al.*, *Biochimica et Biophysica Acta* 953:48-60 (1988); Aarnes *et al.*, *Phytochemistry* 13:2717-2724 (1974); Lea *et al.*, *FEBS Letters* 98:165-168 (1979); Matthews *et al.*, *Canadian Journal of Botany* 57:299-304 (1979), all of which references are incorporated herein in their entirety). The isoenzymes have been found to differ with respect to sensitivity to threonine-mediated feedback inhibition, with both sensitive and insensitive forms being isolated from maize suspension cultures and seedlings (Miflin *et al.*, In: *Nitrogen Assimilation of Plants*, Hewitt *et al.*, (eds.), Academic Press, New York, 335 (1997); Bryan, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York, 5:403 (1980)).

There is evidence that plants also possess a bifunctional enzyme with both aspartate kinase and homoserine dehydrogenase activities (Lea *et al.*, In: *The Chemistry and Biochemistry of Amino Acids*, Barrett *et al.* (eds), London, 5:197 (1985), the entirety of which is herein incorporated by reference; Bryan, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York, 5:161 (1990), the entirety of which is herein incorporated by reference). Clones of these bifunctional enzymes have been isolated from *Arabidopsis thaliana* (Giovanelli *et al.*, In: *The Biochemistry of Plants*, Miflin (ed.), Academic Press, New York 453 (1990), the entirety of which is herein incorporated by reference) carrot (Giovanelli *et al.*, *Plant Physiology* 90:1584-1599 (1989), the entirety of which is herein incorporated by reference), maize (Singh *et al.*, *Amino Acids* 7:165-168 (1994), the entirety of which is herein incorporated by reference) and soybean (Matthews *et al.*, In: *Biosynthesis and Molecular Regulation of Amino Acids in Plants*, p



294, Singh *et al.* (eds.), American Society of Plant Physiologists, Rockville, MD (1992), the entirety of which is herein incorporated by reference).

The next reported enzymatic step leading to methionine biosynthesis in higher plants is the final common reaction shared by other amino acid end products (threonine and isoleucine). The reaction is catalyzed by homoserine kinase and it generates *O*-phosphohomoserine from homoserine, with ATP serving as the phosphate donor. Exceptions are *Pisum sativum* and *Lathyrus sitivus* which synthesize *O*-acetylhomoserine and *O*-oxalylhomoserine, respectively (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997), the entirety of which is herein incorporated by reference). Enteric bacteria use *O*-succinylhomoserine, while several gram-positive bacteria, yeasts and fungi use *O*-acetylhomoserine (formed using homoserine *O*-acetyltransferase (EC 2.3.1.31) (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)). Homoserine kinase has been reported from multiple higher plant sources (Galili, *The Plant Cell* 7:899-906 (1995), the entirety of which is herein incorporated by reference; Rees *et al.*, *Biochemical Journal* 309:999-1107 (1995), the entirety of which is herein incorporated by reference; Bryan *et al.*, *Biochemistry and Biophysics Research Communications* 41:1211-1217 (1970), the entirety of which is herein incorporated by reference; Gengenbach *et al.*, *Crop Science* 18:472-476 (1978), Dotson *et al.*, *Plant Physiology* 91:1602-1608 (1989), the entirety of which is herein incorporated by reference; Dotson *et al.*, *Plant Physiology* 93:98-104 (1989), the entirety of which is herein incorporated by reference). Homoserine kinase isolated from barley and wheat has not been reported to exhibit aspartate-derived amino acid feedback inhibition (Gengenbach *et al.*, *Crop Science* 18:472-476 (1978); Dotson *et al.*, *Plant Physiology* 93:98-104 (1989)). It has been reported that homoserine kinase exhibits feedback regulation in the dicots, pea (Rees *et al.*, *Biochemical Journal* 309:999-1007 (1995), the entirety of which is herein

incorporated by reference) and radish (Bryan *et al.*, *Biochemistry and Biophysics Research Communications* 41:1211-1217 (1970)). Bacterial and yeast homologues have been reported (Azevedo *et al.*, *Phytochemistry* 31:3725-3730 (1992); Azevedo *et al.*, *Phytochemistry* 31:3731-3734 (1992); Brennecke *et al.*, *Phytochemistry* 41:707 (1996); Aarnes, *Plant Science Letters* 9:137-145 (1977)).

Sulfur, in yeast, is incorporated into *O*-acetylhomoserine resulting in homocysteine. This reaction is catalyzed by the *O*-acetylhomoserine sulfhydrylase (EC 4.2.99.10) (also known as *O*-acethomoserine (thiol)-lyase). *O*-acetylhomoserine sulfhydrylase has been reported to be a homotetramer with a molecular weight of 200,000. *O*-acetylhomoserine sulfhydrylase has also been reported to bind four molecules of pyridoxal phosphate (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

In higher plants, the sulfur atom from cysteine and the carbon backbone derived from aspartate used to synthesize methionine are reported to be catalyzed by pyridoxal 5'-phosphate (PLP) dependent enzymes (Ravanel *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 95:7805-7812 (1998), the entirety of which is herein incorporated by reference). The amino acid composition of the *O*-acetylhomoserine sulfhydrylase has also been reported to share sequence similarities to the *E. coli* cystathionine  $\gamma$ -synthase and cystathionine  $\beta$ -lyase and cystathionine  $\gamma$ -lyase from *Saccharomyces cerevisiae* and rats. All of these enzymes thus appear to belong to one protein family, whose members have evolved from an ancestral pyridoxal phosphate enzyme (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

In yeast, the synthesis of cysteine from homocysteine has been reported to require two successive steps,  $\beta$  addition and  $\gamma$  elimination. Cystathionine  $\beta$ -synthase (EC 4.2.1.22) has been reported to catalyze the first reaction where homocysteine and serine yield cystathionine. In *S.*

*cervisiae*, cystathionine  $\beta$ -synthase is encoded by STR4. STR4 encodes a polypeptide of 506 residues which shows extensive sequence similarity to its functional analog in rats. The rat analog has been reported to contain an additional amino-terminal extension of 60 residues. Moreover, the two enzymes have been reported to be closely related to the cyteine synthase from enteric bacteria and plants (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

Cystathionine  $\gamma$ -lyase (EC 4.4.1.1) catalyzes the  $\gamma$  cleavage of cystationine in yeast, the second reported step of the biosynthesis of cysteine from homocysteine. Cystathionine  $\gamma$ -lyase has been reported to have a molecular weight of about 194,000kd. In *S. cerevisiae*, cystathionine  $\gamma$ -lyase is encoded by STR1. A mutation in the *S. cerevisiae* cystathionine  $\gamma$ -lyase gene leads to a nutritional requirement for cysteine or glutathione. The yeast cystathionine  $\gamma$ -lyase belongs to a protein family which includes a functional analog in rats, a Met25p from yeast and cystathionin  $\beta$ -lyase and cystathionin  $\gamma$ -synthase from *E. coli* (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)).

Cystathionine  $\gamma$ -synthase (also known as *O*-succinylhomoserine (thio)-lyase, E.C. 4.2.99.9) catalyzes the first reported reaction which is unique to methionine biosynthesis, thereby committing aspartate pathway flux toward this amino acid. In this reaction, *O*-phosphohomoserine and cysteine serve as substrates for the production of cystathionine. Cystathionine  $\gamma$ -synthase has not been reported to be regulated by aspartate-derived amino acids feedback inhibition (Bright *et al.*, *Biochemical Genetics* 20:229-243 (1982); Arruda *et al.*, *Plant Physiology* 76:442-446 (1984)). Cystathionine  $\gamma$ -synthase has however, been reported to be sensitive to product inhibition by orthophosphate (Lea *et al.*, *Barley: Genetics, Molecular*

*Biology and Biotechnology*, Shewrey (ed.), CAB International, Oxford, 181 (1992); Davies *et al.*, *Plant Science Letters* 9:323-332 (1977)). Cloned cystathionine  $\gamma$ -synthase have been reported from *Arabidopsis thaliana* (Davies *et al.*, *Plant Physiology* 62:536-541 (1978)). It has been reported that methionine levels are modulated via regulation of cystathionine-synthase (Matthews *et al.*, *Zeitschrift für Naturforschung, Section Bioscience* 34:1177-1185 (1979-2724 (1974); Lea *et al.*, *FEBS Letters* 98:165 (1979), all of which references are incorporated herein in their entirety).

Cystathionine  $\beta$ -lyase catalyzes the next reaction in the biosynthesis of methionine. This reaction generates homocysteine, pyruvate and ammonia from the enzymatic decomposition of cystathionine. Evidence for isoenzymes which differ with respect to cellular localization have been reported for barley (Matthews *et al.*, *Canadian Journal of Botany* 57:299-304 (1979)) and spinach (Rognes *et al.*, *Nature* 287:357-359 (1980), the entirety of which is herein incorporated by reference).

De novo synthesis of methionine from homocysteine uses a methyl group which originates from single-carbon metabolism. In this metabolism, derivatives of tetrahydrofolate transfer one-carbon groups at the oxidation levels of methanol, formaldehyde and formate to acceptor molecules. Single-carbon derivatives of tetrahydrofolate are required for the biosynthesis of methionine, purine nucleotides and thymidylate as well as for the synthesis of N-formylmethionine in the mitochondrion. *S. cerevisiae* possesses two complete sets of folate interconversion enzymes, one located in the cytosol (methionyl-tRNA synthetase, EC 6.1.1.10) and the other located in the mitochondrion (methionyl t-RNA synthetase, EC 6.1.1.10) (Thomas and Surdin-Kerjan, *Microbiol. Mol. Biol. Rev.* 61:503-532 (1997)) and in plants including the

chloroplast (Menand *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)*, 95:11014-11019 (1998), the entirety of which is herein incorporated by reference).

Methionine synthase generates methionine from homocysteine by a methylation reaction and thus represents the final step of the methionine biosynthetic pathway. Methionine synthase is also sometimes referred to as 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase. N-methyltetrahydrofolate serves as the methyl donor in this reaction, which occurs in the absence of cobalamin (Giovanelli *et al.*, *Plant Physiology* 90:1577-1583 (1989), the entirety of which is herein incorporated by reference; Green *et al.*, *Crop Science* 14:827-830 (1974), the entirety of which is herein incorporated by reference).

## II. METHIONINE DEGRADATION PATHWAY

Plants contain a pathway for the degradation of L-methionine. This degradation pathway includes the following enzymes: methionine adenosyltransferase (EC 2.5.1.6), methionine S-methyltransferase (EC 2.1.1.12), adenosylmethionine hydrolase (EC 3.3.1.2), homocysteine S-methyltransferase (EC 2.1.1.10) and S-adenosyl-methionine decarboxylase (EC 4.1.1.50).

The reported first step in the catabolism of methionine is the ATP-dependent conversion to S-adenosylmethionine (AdoMet), which is catalyzed by the enzyme methionine adenosyltransferase, also known as S-adenosylmethionine synthetase. Methionine adenosyltransferase enzyme has been characterized from several plant sources (Aarnes, *Plant Science Letters* 10:381 (1977), the entirety of which is herein incorporated by reference; Mathur *et al.*, *Biochimica and Biophysica Acta* 1078:161-170 (1991), the entirety of which is herein incorporated by reference; Kim *et al.*, *Journal of Biochemical and Molecular Biology* 28:100 (1995), the entirety of which is herein incorporated by reference) and nucleic acid molecules (genomic and cDNA) have also been obtained from a variety of sources (Izhaki *et al.*, *Plant*

*Physiology* 108:841-842 (1995), the entirety of which is herein incorporated by reference; Espartero *et al.*, *Molecular Biology Plant* 25:217-237 (1994), the entirety of which is herein incorporated by reference). Regulation of methionine adenosyltransferase activity has been observed for the enzyme from *Glycine max* (soybean). In *Glycine max*, methionine adenosyltransferase was reportedly inhibited by S-adenosylmethionine (Kim *et al.*, *Journal of Biochemical and Molecular Biology* 28:100 (1995). Studies have also reported that the levels of methionine adenosyltransferase appear to fluctuate in response to hormonal or environmental conditions such as gibberellic acid (Mathur *et al.*, *Biochimica and Biophysica Acta* 1162:289-290 (1993), the entirety of which is herein incorporated by reference; Mathur *et al.*, *Biochimica and Biophysica Acta* 1137:338-348 (1992), the entirety of which is herein incorporated by reference), salt stress (Espartero *et al.*, *Molecular Biology Plant* 25:217-227 (1994) the entirety of which is herein incorporated by reference) and wounding (Kim *et al.*, *Plant Cell Reports* 13:340 (1994), the entirety of which is herein incorporated by reference). It has also been reported that methionine adenosyltransferase may play a role in the lignification process (Peleman *et al.*, *Plant Cell* 1:81 (1989), the entirety of which is herein incorporated by reference).

AdoMet is further catabolized by several enzymes and has been reported to serve a variety of metabolic functions including that of a methyl donor (Cossins, *The Biochemistry of Plants* 11:317 Devis (ed.), Academic Press, San Diego (1987), the entirety of which is herein incorporated by reference) that of a precursor for polyamine biosynthesis (Tiburico *et al.*, *The Biochemistry of Plants* 16:283 (1990), the entirety of which is herein incorporated by reference) and that of a precursor for ethylene biosynthesis (Kende, *Plant Physiology* 91:1-4 (1989), the entirety of which is herein incorporated by reference; Flurh *et al.*, *Critical Review of Plant*

*Science* 15:479 (1996), the entirety of which is herein incorporated by reference). In each case, enzymes are present to regenerate methionine from the sulfur-containing backbone resulting in no net loss of methionine.

An enzyme involved in AdoMet catabolism is adenosylmethionine hydrolase (EC 3.3.1.2) which converts AdoMet to methylthioadenosine and L-homoserine. L-homoserine is further metabolized during the biosynthesis of polyamines and ethylene and methylthioadenosine is recycled to methionine. In yeast, a form of adenosylmethionine hydrolase (EC 3.1.1.1) has been reported (<http://www.ncbi.nlm.nih.gov/htbin-post/Entrez/query> (1998)).

Another enzyme for which AdoMet is a substrate for is homocysteine S-methyltransferase. Homocysteine S-methyltransferase catalyzes the combination of AdoMet, with L-homocysteine to produce both S-adenosyl-L-homocysteine and L-methionine. Another enzyme has been described which generates S-adenosyl-L-homocysteine from AdoMet. This enzyme is called methionine S-methyltransferase and it catalyzes the reaction in which S-adenosyl-L-homocysteine reacts with L-methionine to generate S-adenosyl-L-homocysteine and S-methyl-L-methionine. AdoMet can also be decarboxylated by adenosyl methionine decarboxylase, which generates (5-deoxy-5-adenosyl) (3-aminopropyl) methylsulfonium salt.

### **III. EXPRESSED SEQUENCE TAG NUCLEIC ACID MOLECULES**

Expressed sequence tags, or ESTs are randomly sequenced members of a cDNA library (or complementary DNA)(McCombie *et al.*, *Nature Genetics* 1:124-130 (1992); Kurata *et al.*, *Nature Genetics* 8:365-372 (1994); Okubo *et al.*, *Nature Genetics* 2:173-179 (1992), all of which references are incorporated herein in their entirety). The randomly selected clones comprise insets that can represent a copy of up to the full length of a mRNA transcript.

Using conventional methodologies, cDNA libraries can be constructed from the mRNA (messenger RNA) of a given tissue or organism using poly dT primers and reverse transcriptase (Efstratiadis *et al.*, *Cell* 7:279-3680 (1976), the entirety of which is herein incorporated by reference; Higuchi *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 73:3146-3150 (1976), the entirety of which is herein incorporated by reference; Maniatis *et al.*, *Cell* 8:163-182 (1976) the entirety of which is herein incorporated by reference; Land *et al.*, *Nucleic Acids Res.* 9:2251-2266 (1981), the entirety of which is herein incorporated by reference; Okayama *et al.*, *Mol. Cell. Biol.* 2:161-170 (1982), the entirety of which is herein incorporated by reference; Gubler *et al.*, *Gene* 25:263-269 (1983), the entirety of which is herein incorporated by reference).

Several methods may be employed to obtain full-length cDNA constructs. For example, terminal transferase can be used to add homopolymeric tails of dC residues to the free 3' hydroxyl groups (Land *et al.*, *Nucleic Acids Res.* 9:2251-2266 (1981), the entirety of which is herein incorporated by reference). This tail can then be hybridized by a poly dG oligo which can act as a primer for the synthesis of full length second strand cDNA. Okayama and Berg, *Mol. Cell. Biol.* 2:161-170 (1982), the entirety of which is herein incorporated by reference, report a method for obtaining full length cDNA constructs. This method has been simplified by using synthetic primer-adapters that have both homopolymeric tails for priming the synthesis of the first and second strands and restriction sites for cloning into plasmids (Coleclough *et al.*, *Gene* 34:305-314 (1985), the entirety of which is herein incorporated by reference) and bacteriophage vectors (Krawinkel *et al.*, *Nucleic Acids Res.* 14:1913 (1986), the entirety of which is herein incorporated by reference; Han *et al.*, *Nucleic Acids Res.* 15:6304 (1987), the entirety of which is herein incorporated by reference).



These strategies have been coupled with additional strategies for isolating rare mRNA populations. For example, a typical mammalian cell contains between 10,000 and 30,000 different mRNA sequences (Davidson, *Gene Activity in Early Development*, 2nd ed., Academic Press, New York (1976), the entirety of which is herein incorporated by reference). The number of clones required to achieve a given probability that a low-abundance mRNA will be present in a cDNA library is  $N = (\ln(1-P))/(\ln(1-1/n))$  where N is the number of clones required, P is the probability desired and 1/n is the fractional proportion of the total mRNA that is represented by a single rare mRNA (Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, 2nd ed., Cold Spring Harbor Laboratory Press (1989), the entirety of which is herein incorporated by reference).

A method to enrich preparations of mRNA for sequences of interest is to fractionate by size. One such method is to fractionate by electrophoresis through an agarose gel (Pennica *et al.*, *Nature* 301:214-221 (1983), the entirety of which is herein incorporated by reference). Another such method employs sucrose gradient centrifugation in the presence of an agent, such as methylmercuric hydroxide, that denatures secondary structure in RNA (Schweinfest *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 79:4997-5000 (1982), the entirety of which is herein incorporated by reference).

A frequently adopted method is to construct equalized or normalized cDNA libraries (Ko, *Nucleic Acids Res.* 18:5705-5711 (1990), the entirety of which is herein incorporated by reference; Patanjali *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:1943-1947 (1991), the entirety of which is herein incorporated by reference). Typically, the cDNA population is normalized by subtractive hybridization (Schmid *et al.*, *J. Neurochem.* 48:307-312 (1987), the entirety of which is herein incorporated by reference; Fargnoli *et al.*, *Anal. Biochem.* 187:364-373 (1990), the

entirety of which is herein incorporated by reference; Travis *et al.*, *Proc. Natl. Acad. Sci (U.S.A.)* 85:1696-1700 (1988), the entirety of which is herein incorporated by reference; Kato, *Eur. J. Neurosci.* 2:704-711 (1990); and Schweinfest *et al.*, *Genet. Anal. Tech. Appl.* 7:64-70 (1990), the entirety of which is herein incorporated by reference). Subtraction represents another method for reducing the population of certain sequences in the cDNA library (Swaroop *et al.*, *Nucleic Acids Res.* 19:1954 (1991), the entirety of which is herein incorporated by reference).

ESTs can be sequenced by a number of methods. Two basic methods may be used for DNA sequencing, the chain termination method of Sanger *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 74:5463-5467 (1977), the entirety of which is herein incorporated by reference and the chemical degradation method of Maxam and Gilbert, *Proc. Nat. Acad. Sci. (U.S.A.)* 74:560-564 (1977), the entirety of which is herein incorporated by reference. Automation and advances in technology such as the replacement of radioisotopes with fluorescence-based sequencing have reduced the effort required to sequence DNA (Craxton, *Methods* 2:20-26 (1991), the entirety of which is herein incorporated by reference; Ju *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 92:4347-4351 (1995), the entirety of which is herein incorporated by reference; Tabor and Richardson, *Proc. Natl. Acad. Sci. (U.S.A.)* 92:6339-6343 (1995), the entirety of which is herein incorporated by reference). Automated sequencers are available from, for example, Pharmacia Biotech, Inc., Piscataway, New Jersey (Pharmacia ALF), LI-COR, Inc., Lincoln, Nebraska (LI-COR 4,000) and Millipore, Bedford, Massachusetts (Millipore BaseStation).

In addition, advances in capillary gel electrophoresis have also reduced the effort required to sequence DNA and such advances provide a rapid high resolution approach for sequencing DNA samples (Swerdlow and Gesteland, *Nucleic Acids Res.* 18:1415-1419 (1990); Smith, *Nature* 349:812-813 (1991); Luckey *et al.*, *Methods Enzymol.* 218:154-172 (1993); Lu *et al.*, *J.*

*Chromatog. A.* 680:497-501 (1994); Carson *et al.*, *Anal. Chem.* 65:3219-3226 (1993); Huang *et al.*, *Anal. Chem.* 64:2149-2154 (1992); Kheterpal *et al.*, *Electrophoresis* 17:1852-1859 (1996); Quesada and Zhang, *Electrophoresis* 17:1841-1851 (1996); Baba, *Yakugaku Zasshi* 117:265-281 (1997), all of which are herein incorporated by reference in their entirety).

ESTs longer than 150 nucleotides have been found to be useful for similarity searches and mapping (Adams *et al.*, *Science* 252:1651-1656 (1991), herein incorporated by reference). ESTs, which can represent copies of up to the full length transcript, may be partially or completely sequenced. Between 150-450 nucleotides of sequence information is usually generated as this is the length of sequence information that is routinely and reliably produced using single run sequence data. Typically, only single run sequence data is obtained from the cDNA library (Adams *et al.*, *Science* 252:1651-1656 (1991). Automated single run sequencing typically results in an approximately 2-3% error or base ambiguity rate (Boguski *et al.*, *Nature Genetics* 4:332-333 (1993), the entirety of which is herein incorporated by reference).

EST databases have been constructed or partially constructed from, for example, *C. elegans* (McCombie *et al.*, *Nature Genetics* 1:124-131 (1992)), human liver cell line HepG2 (Okubo *et al.*, *Nature Genetics* 2:173-179 (1992)), human brain RNA (Adams *et al.*, *Science* 252:1651-1656 (1991); Adams *et al.*, *Nature* 355:632-635 (1992)), *Arabidopsis*, (Newman *et al.*, *Plant Physiol.* 106:1241-1255 (1994)); and rice (Kurata *et al.*, *Nature Genetics* 8:365-372 (1994)).

#### IV. SEQUENCE COMPARISONS

A characteristic feature of a DNA sequence is that it can be compared with other DNA sequences. Sequence comparisons can be undertaken by determining the similarity of the test or query sequence with sequences in publicly available or proprietary databases ("similarity

analysis”) or by searching for certain motifs (“intrinsic sequence analysis”)(e.g. *cis* elements)(Coulson, *Trends in Biotechnology* 12:76-80 (1994), the entirety of which is herein incorporated by reference); Birren *et al.*, *Genome Analysis 1*: Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York 543-559 (1997), the entirety of which is herein incorporated by reference).

Similarity analysis includes database search and alignment. Examples of public databases include the DNA Database of Japan (DDBJ)(<http://www.ddbj.nig.ac.jp/>); Genebank (<http://www.ncbi.nlm.nih.gov/Web/Search/Index.html>); and the European Molecular Biology Laboratory Nucleic Acid Sequence Database (EMBL) ([http://www.ebi.ac.uk/ebi\\_docs/embl\\_db/embl-db.html](http://www.ebi.ac.uk/ebi_docs/embl_db/embl-db.html)). Other appropriate databases include dbEST (<http://www.ncbi.nlm.nih.gov/dbEST/index.html>), SwissProt ([http://www.ebi.ac.uk/ebi\\_docs/swisprot\\_db/swisshome.html](http://www.ebi.ac.uk/ebi_docs/swisprot_db/swisshome.html)), PIR (<http://www-nbrt.georgetown.edu/pir/>) and The Institute for Genome Research (<http://www.tigr.org/tdb/tdb.html>)

A number of different search algorithms have been developed, one example of which are the suite of programs referred to as BLAST programs. There are five implementations of BLAST, three designed for nucleotide sequences queries (BLASTN, BLASTX and TBLASTX) and two designed for protein sequence queries (BLASTP and TBLASTN) (Coulson, *Trends in Biotechnology* 12:76-80 (1994); Birren *et al.*, *Genome Analysis 1*, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York 543-559 (1997)).

BLASTN takes a nucleotide sequence (the query sequence) and its reverse complement and searches them against a nucleotide sequence database. BLASTN was designed for speed, not maximum sensitivity and may not find distantly related coding sequences. BLASTX takes a

nucleotide sequence, translates it in three forward reading frames and three reverse complement reading frames and then compares the six translations against a protein sequence database.

BLASTX is useful for sensitive analysis of preliminary (single-pass) sequence data and is tolerant of sequencing errors (Gish and States, *Nature Genetics* 3:266-272 (1993), the entirety of which is herein incorporated by reference). BLASTN and BLASTX may be used in concert for analyzing EST data (Coulson, *Trends in Biotechnology* 12:76-80 (1994); Birren *et al.*, *Genome Analysis* 1:543-559 (1997)).

Given a coding nucleotide sequence and the protein it encodes, it is often preferable to use the protein as the query sequence to search a database because of the greatly increased sensitivity to detect more subtle relationships. This is due to the larger alphabet of proteins (20 amino acids) compared with the alphabet of nucleic acid sequences (4 bases), where it is far easier to obtain a match by chance. In addition, with nucleotide alignments, only a match (positive score) or a mismatch (negative score) is obtained, but with proteins, the presence of conservative amino acid substitutions can be taken into account. Here, a mismatch may yield a positive score if the non-identical residue has physical/chemical properties similar to the one it replaced. Various scoring matrices are used to supply the substitution scores of all possible amino acid pairs. A general purpose scoring system is the BLOSUM62 matrix (Henikoff and Henikoff, *Proteins* 17:49-61 (1993), the entirety of which is herein incorporated by reference), which is currently the default choice for BLAST programs. BLOSUM62 is tailored for alignments of moderately diverged sequences and thus may not yield the best results under all conditions. Altschul, *J. Mol. Biol.* 36:290-300 (1993), the entirety of which is herein incorporated by reference, describes a combination of three matrices to cover all contingencies. This may improve sensitivity, but at the expense of slower searches. In practice, a single

BLOSUM62 matrix is often used but others (PAM40 and PAM250) may be attempted when additional analysis is necessary. Low PAM matrices are directed at detecting very strong but localized sequence similarities, whereas high PAM matrices are directed at detecting long but weak alignments between very distantly related sequences.

Homologues in other organisms are available that can be used for comparative sequence analysis. Multiple alignments are performed to study similarities and differences in a group of related sequences. CLUSTAL W is a multiple sequence alignment package that performs progressive multiple sequence alignments based on the method of Feng and Doolittle, *J. Mol. Evol.* 25:351-360 (1987), the entirety of which is herein incorporated by reference. Each pair of sequences is aligned and the distance between each pair is calculated; from this distance matrix, a guide tree is calculated and all of the sequences are progressively aligned based on this tree. A feature of the program is its sensitivity to the effect of gaps on the alignment; gap penalties are varied to encourage the insertion of gaps in probable loop regions instead of in the middle of structured regions. Users can specify gap penalties, choose between a number of scoring matrices, or supply their own scoring matrix for both pairwise alignments and multiple alignments. CLUSTAL W for UNIX and VMS systems is available at: <ftp://ebi.ac.uk>. Another program is MACAW (Schuler *et al.*, *Proteins Struct. Func. Genet.* 9:180-190 (1991), the entirety of which is herein incorporated by reference, for which both Macintosh and Microsoft Windows versions are available. MACAW uses a graphical interface, provides a choice of several alignment algorithms and is available by anonymous ftp at: [ncbi.nlm.nih.gov](ftp://ncbi.nlm.nih.gov) (directory/pub/macaw).

Sequence motifs are derived from multiple alignments and can be used to examine individual sequences or an entire database for subtle patterns. With motifs, it is sometimes

possible to detect distant relationships that may not be demonstrable based on comparisons of primary sequences alone. Currently, the largest collection of sequence motifs in the world is PROSITE (Bairoch and Bucher, *Nucleic Acid Research* 22:3583-3589 (1994), the entirety of which is herein incorporated by reference). PROSITE may be accessed via either the ExPASy server on the World Wide Web or anonymous ftp site. Many commercial sequence analysis packages also provide search programs that use PROSITE data.

A resource for searching protein motifs is the BLOCKS E-mail server developed by Henikoff, *Trends Biochem Sci.* 18:267-268 (1993), the entirety of which is herein incorporated by reference; Henikoff and Henikoff, *Nucleic Acid Research* 19:6565-6572 (1991), the entirety of which is herein incorporated by reference; Henikoff and Henikoff, *Proteins* 17:49-61 (1993). BLOCKS searches a protein or nucleotide sequence against a database of protein motifs or “blocks.” Blocks are defined as short, ungapped multiple alignments that represent highly conserved protein patterns. The blocks themselves are derived from entries in PROSITE as well as other sources. Either a protein query or a nucleotide query can be submitted to the BLOCKS server; if a nucleotide sequence is submitted, the sequence is translated in all six reading frames and motifs are sought for these conceptual translations. Once the search is completed, the server will return a ranked list of significant matches, along with an alignment of the query sequence to the matched BLOCKS entries.

Conserved protein domains can be represented by two-dimensional matrices, which measure either the frequency or probability of the occurrences of each amino acid residue and deletions or insertions in each position of the domain. This type of model, when used to search against protein databases, is sensitive and usually yields more accurate results than simple motif searches. Two popular implementations of this approach are profile searches such as GCG

program ProfileSearch and Hidden Markov Models (HMMs)(Krough *et al.*, *J. Mol. Biol.* 235:1501-1531, (1994); Eddy, *Current Opinion in Structural Biology* 6:361-365, (1996), both of which are herein incorporated by reference in their entirety). In both cases, a large number of common protein domains have been converted into profiles, as present in the PROSITE library, or HMM models, as in the Pfam protein domain library (Sonnhammer *et al.*, *Proteins* 28:405-420 (1997), the entirety of which is herein incorporated by reference). Pfam contains more than 500 HMM models for enzymes, transcription factors, signal transduction molecules and structural proteins. Protein databases can be queried with these profiles or HMM models, which will identify proteins containing the domain of interest. For example, HMMSW or HMMFS, two programs in a public domain package called HMMER (Sonnhammer *et al.*, *Proteins* 28:405-420 (1997)) can be used.

PROSITE and BLOCKS represent collected families of protein motifs. Thus, searching these databases entails submitting a single sequence to determine whether or not that sequence is similar to the members of an established family. Programs working in the opposite direction compare a collection of sequences with individual entries in the protein databases. An example of such a program is the Motif Search Tool, or MoST (Tatusov *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:12091-12095 (1994), the entirety of which is herein incorporated by reference). On the basis of an aligned set of input sequences, a weight matrix is calculated by using one of four methods (selected by the user). A weight matrix is simply a representation, position by position of how likely a particular amino acid will appear. The calculated weight matrix is then used to search the databases. To increase sensitivity, newly found sequences are added to the original data set, the weight matrix is recalculated and the search is performed again. This procedure continues until no new sequences are found.



## SUMMARY OF THE INVENTION

The present invention provides a substantially purified nucleic acid molecule that encodes a maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of: (a) methionine adenosyltransferase, (b) S-adenosyl-methionine decarboxylase, (c) aspartate kinase, (d) aspartate-semialdehyde dehydrogenase, (e) cystathionine gamma-synthase, (f) cystathionine beta-lyase, and (g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase.

The present invention also provides a substantially purified nucleic acid molecule that encodes a plant methionine pathway enzyme or fragment thereof, wherein the nucleic acid molecule is selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a

maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof.

A substantially purified maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of (a) methionine adenosyltransferase or fragment thereof; (b) S-adenosyl-methionine decarboxylase or fragment thereof; (c) aspartate kinase or fragment thereof; (d) aspartate-semialdehyde dehydrogenase or fragment thereof; (e) cystathionine gamma-synthase or fragment thereof; (f) cystathionine beta-lysase or fragment thereof; and (g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof.

The present invention also provides a substantially purified maize or soybean methionine pathway protein or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1 through SEQ ID NO: 3204.

The present invention also provides a substantially purified maize or soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479.

The present invention also provides a substantially purified maize or soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479.

The present invention also provides a substantially purified maize or soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623.

The present invention also provides a substantially purified maize or soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623.

The present invention also provides a substantially purified maize or soybean aspartate kinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648.

The present invention also provides a substantially purified maize or soybean aspartate kinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648.

The present invention also provides a substantially purified maize or soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a

complement of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654.

The present invention also provides a substantially purified maize or soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654.

The present invention also provides a substantially purified maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660.

The present invention also provides a substantially purified maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660.

The present invention also provides a substantially purified maize or soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665.

The present invention also provides a substantially purified maize or soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected

from the group consisting of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665.

The present invention also provides a substantially purified maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992.

The present invention also provides a substantially purified maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992.

The present invention also provides a substantially purified maize or adenosylhomocysteinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199.

The present invention also provides a substantially purified maize or adenosylhomocysteinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199.

The present invention also provides a substantially purified maize or cystathionine  $\beta$ -synthase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1631 through SEQ ID NO: 1632.

The present invention also provides a substantially purified maize or cystathionine  $\beta$ -synthase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1631 through SEQ ID NO: 1632.

The present invention also provides a substantially purified maize or cystathionine  $\gamma$ -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204.

The present invention also provides a substantially purified maize or cystathionine  $\gamma$ -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204.

The present invention also provides a substantially purified maize or *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 3200 through SEQ ID NO: 3202.

The present invention also provides a substantially purified maize or *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 3200 through SEQ ID NO: 3202.

The present invention also provides a purified antibody or fragment thereof which is capable of specifically binding to a specific maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme or fragment thereof is encoded by a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479 or a substantially purified maize or soybean methionine adenosyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ

ID NO: 2623 or a substantially purified maize or soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean aspartate kinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648 or a substantially purified maize or soybean aspartate kinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654 or a substantially purified maize or soybean enzyme aspartate-semialdehyde dehydrogenase or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654.



The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or a substantially purified maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665 or a substantially purified maize or soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2661 through SEQ ID NO: 2665.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic

acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992 or a substantially purified maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199 or a substantially purified maize or soybean adenosylhomocysteinase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1631 through SEQ ID NO: 1632 or a substantially purified maize or soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1631 through SEQ ID NO: 1632.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204 or a substantially purified maize or soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204.

The present invention also provides a substantially purified antibody or fragment thereof, the antibody or fragment thereof capable of specifically binding to a maize or a soybean *O*-acetylhomoserine enzyme or fragment thereof encoded by a first nucleic acid molecule which specifically hybridizes to a second nucleic acid molecule, the second nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a complement of SEQ ID NO: 3200 through SEQ ID NO: 3202 or a substantially purified maize or soybean *O*-acetylhomoserine enzyme or fragment thereof encoded by a nucleic acid sequence selected from the group consisting of SEQ ID NO: 3200 through SEQ ID NO: 3202.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; (B) a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of (a) a nucleic acid sequence which encodes for methionine adenosyltransferase or fragment thereof; (b) a nucleic acid sequence which encodes for S-adenosyl-methionine decarboxylase or fragment thereof; (c) a nucleic acid

sequence which encodes for aspartate kinase or fragment thereof; (d) a nucleic acid sequence which encodes for aspartate-semialdehyde dehydrogenase or fragment thereof; (e) a nucleic acid sequence which encodes for cystathionine gamma-synthase or a fragment thereof; (f) a nucleic acid sequence which encodes for cystathionine beta-lyase or a fragment thereof; (g) a nucleic acid sequence which encodes for 5-methyltetrahydropteroyl- triglutamate-homocysteine-S-methyltransferase or a fragment thereof; and (h) a nucleic acid sequence which is complementary to any of the nucleic acid sequences of (a) through (g); and (C) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to (B) a structural nucleic acid molecule, wherein the structural nucleic acid molecule encodes a plant methionine pathway enzyme or fragment thereof, the structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof; which is linked to (C) a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to (B) a structural nucleic acid molecule, wherein the structural nucleic acid molecule is selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or

fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; which is linked to (C) a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to (B) a transcribed nucleic acid molecule with a transcribed strand and a non-transcribed strand, wherein the transcribed strand is complementary to a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof; which is

linked to (C) a 3' non-translated sequence that functions in plant cells to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a transformed plant having a nucleic acid molecule which comprises: (A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule; which is linked to: (B) a transcribed nucleic acid molecule with a transcribed strand and a non-transcribed strand, wherein a transcribed mRNA of the transcribed strand is complementary to an endogenous mRNA molecule having a nucleic acid sequence selected from the group consisting of an endogenous mRNA molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and an endogenous mRNA molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; which is linked to (C) a 3'

non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of the mRNA molecule.

The present invention also provides a method for determining a level or pattern in a plant cell of an enzyme in a plant metabolic pathway comprising: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence of SEQ ID NO: 1 through SEQ ID NO: 3204 or compliments thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said enzyme; (B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and (C) detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said enzyme in said plant metabolic pathway.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue comprising: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complement thereof or fragment of either, with a complementary nucleic acid molecule obtained from the plant cell or plant tissue, wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue permits the detection of the

plant methionine pathway enzyme; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue; and (C) detecting the level or pattern of the complementary nucleic acid, wherein the detection of the complementary nucleic acid is predictive of the level or pattern of the plant methionine pathway enzyme.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue comprising: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule comprising a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a



soybean cytsathionine  $\gamma$ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, with a complementary nucleic acid molecule obtained from the plant cell or plant tissue, wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue permits the detection of the plant methionine pathway enzyme; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue; and (C) detecting the level or pattern of the complementary nucleic acid, wherein the detection of the complementary nucleic acid is predictive of the level or pattern of the plant methionine pathway enzyme.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue under evaluation which comprises assaying the concentration of a molecule, whose concentration is dependent upon the expression of a gene, the gene specifically hybridizes to a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof, in comparison to the concentration of that molecule present in a reference plant cell or a reference plant tissue with a known level or pattern of the plant methionine pathway enzyme, wherein the assayed concentration of the molecule is compared to the assayed concentration of the molecule in the reference plant cell or reference plant tissue with the known level or pattern of the plant methionine pathway enzyme.

The present invention also provides a method for determining a level or pattern of a plant methionine pathway enzyme in a plant cell or plant tissue under evaluation which comprises assaying the concentration of a molecule, whose concentration is dependent upon the expression

of a gene, the gene specifically hybridizes to a nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or complement thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof, in comparison to the concentration of that molecule present in a reference plant cell or a reference plant tissue with a known level or pattern of the plant methionine pathway enzyme, wherein the assayed concentration of the molecule is compared to the assayed concentration of the molecule in the reference plant cell or the reference plant tissue with the known level or pattern of the plant methionine pathway enzyme.

A method of determining a mutation in a plant whose presence is predictive of a mutation affecting a level or pattern of a protein comprising the steps: (A) incubating, under conditions

permitting nucleic acid hybridization, a marker nucleic acid, said marker nucleic acid selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having a nucleic acid sequence selected from the group of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof and a complementary nucleic acid molecule obtained from said plant, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting said level or pattern of said plant methionine pathway enzyme in said plant; (B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant; and (C) detecting the presence of said polymorphism, wherein the detection of said polymorphism is predictive of said mutation.

The present invention also provides a method for determining a mutation in a plant whose presence is predictive of a mutation affecting the level or pattern of a plant methionine pathway enzyme comprising the steps: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule comprising a nucleic acid molecule that is linked to a gene, the gene specifically hybridizes to a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof and a complementary nucleic acid molecule obtained from the plant, wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting the level or pattern of the plant methionine pathway enzyme in the plant; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule

obtained from the plant; and (C) detecting the presence of the polymorphism, wherein the detection of the polymorphism is predictive of the mutation.

The present invention also provides a method for determining a mutation in a plant whose presence is predictive of a mutation affecting the level or pattern of a plant methionine pathway enzyme comprising the steps: (A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, the marker nucleic acid molecule comprising a nucleic acid molecule that is linked to a gene, the gene specifically hybridizes to a nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or complement thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or complement thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof and a complementary nucleic acid molecule obtained from the plant,

wherein nucleic acid hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting the level or pattern of the plant methionine pathway enzyme in the plant; (B) permitting hybridization between the marker nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant; and (C) detecting the presence of the polymorphism, wherein the detection of the polymorphism is predictive of the mutation.

A method of producing a plant containing an overexpressed protein comprising: (A) transforming said plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein said promoter region is linked to a structural region, wherein said structural region has a nucleic acid sequence selected from group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 wherein said structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein said functional nucleic acid molecule results in overexpression of the protein; and (B) growing said transformed plant.

The present invention also provides a method of producing a plant containing an overexpressed plant methionine enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof; wherein the structural region is linked to a 3' non-translated sequence that functions in the plant to cause

termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in overexpression of the plant methionine pathway enzyme; and (B) growing the transformed plant.

The present invention also provides a method of producing a plant containing an overexpressed plant methionine pathway enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; wherein the structural region is

linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in overexpression of the plant methionine pathway enzyme protein; and (B) growing the transformed plant.

The present invention also provides a method of producing a plant containing reduced levels of a plant methionine pathway enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204; wherein the structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in co-suppression of the plant methionine pathway enzyme protein; and (B) growing the transformed plant.

The present invention also provides a method of producing a plant containing reduced levels of a plant methionine pathway enzyme comprising: (A) transforming the plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein the promoter region is linked to a structural region, wherein the structural region comprises a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a

nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; wherein the structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and wherein the functional nucleic acid molecule results in co-suppression of the plant methionine pathway enzyme; and (B) growing the transformed plant.

The present invention also provides a method for reducing expression of a plant methionine pathway enzyme in a plant comprising: (A) transforming the plant with a nucleic acid molecule, the nucleic acid molecule having an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule, wherein the exogenous promoter region is linked to a transcribed nucleic acid molecule having a transcribed strand and a non-transcribed strand, wherein the transcribed strand is complementary to a nucleic acid molecule having a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragments of either and the transcribed strand is complementary



to an endogenous mRNA molecule; and wherein the transcribed nucleic acid molecule is linked to a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and (B) growing the transformed plant.

The present invention also provides a method for reducing expression of a plant methionine pathway enzyme in a plant comprising: (A) transforming the plant with a nucleic acid molecule, the nucleic acid molecule having an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule, wherein the exogenous promoter region is linked to a transcribed nucleic acid molecule having a transcribed strand and a non-transcribed strand, wherein a transcribed mRNA of the transcribed strand is complementary to a nucleic acid molecule selected from the group consisting of an endogenous mRNA molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment

thereof, an endogenous mRNA molecule that encodes a maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and an endogenous mRNA molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof; and wherein the transcribed nucleic acid molecule is linked to a 3' non-translated sequence that functions in the plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of a mRNA molecule; and (B) growing the transformed plant.

The present invention also provides a method of determining an association between a polymorphism and a plant trait comprising: (A) hybridizing a nucleic acid molecule specific for the polymorphism to genetic material of a plant, wherein the nucleic acid molecule has a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragment thereof; and (B) calculating the degree of association between the polymorphism and the plant trait.

The present invention also provides a method of determining an association between a polymorphism and a plant trait comprising: (A) hybridizing a nucleic acid molecule specific for the polymorphism to genetic material of a plant, wherein the nucleic acid molecule is selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme complement thereof or fragment of either, a

nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either; a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cytsathionine  $\gamma$ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either and (B) calculating the degree of association between the polymorphism and the plant trait.

The present invention also provides a method of isolating a nucleic acid that encodes a plant methionine pathway enzyme or fragment thereof comprising: (A) incubating under conditions permitting nucleic acid hybridization, a first nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragment of either with a complementary second nucleic acid molecule obtained from a plant cell or plant tissue; (B) permitting hybridization between the first nucleic acid molecule and the second nucleic acid molecule obtained from the plant cell or plant tissue; and (C) isolating the second nucleic acid molecule.

The present invention also provides a method of isolating a nucleic acid molecule that encodes a plant methionine pathway enzyme or fragment thereof comprising: (A) incubating under conditions permitting nucleic acid hybridization, a first nucleic acid molecule selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean

methionine adenosyltransferase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cytsathionine  $\gamma$ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, with a complementary second nucleic acid molecule obtained from a plant cell or plant tissue; (B) permitting hybridization between the plant methionine pathway nucleic acid molecule and the complementary nucleic acid molecule obtained from the plant cell or plant tissue; and (C) isolating the second nucleic acid molecule.

## **DETAILED DESCRIPTION OF THE INVENTION**

### **Definitions and Agents of the Present Invention**

#### **Definitions:**

As used herein, a methionine pathway enzyme is any enzyme that is associated with the synthesis or degradation of methionine.

As used herein, a methionine synthesis enzyme is any enzyme that is associated with the synthesis of methionine.

As used herein, a methionine degradation enzyme is any enzyme that is associated with the degradation of methionine.

As used herein, methionine adenosyltransferase is any enzyme that catalyzes the conversion of methionine to S-adenosylmethionine.

As used herein, S-adenosylmethionine decarboxylase is any enzyme that catalyzes the reaction that converts S-adenosylmethionine to (5-deoxy-5-adenosyl)(3-aminopropyl) methylsulfonium salt.

As used herein, aspartate kinase is any enzyme that catalyzes the conversion of aspartate to  $\beta$ -aspartyl phosphate.

As used herein, aspartate semialdehyde dehydrogenase is any enzyme that catalyzes the conversion of  $\beta$ -aspartyl phosphate to aspartate-semialdehyde via an NADPH-dependent reaction.

As used herein, *O*-succinylhomoserine (thiol)-lyase refers to any enzyme that catalyzes the conversion of *O*-phosphohomoserine to and cysteine to cystathionine.

As used herein, cystathionine  $\beta$ -lyase is any enzyme that catalyzes the conversion of cystathionine to homocysteine, pyruvate and ammonia.

As used herein, 5-methyltetrahydropterolytriglutamate-homocysteine S-methyltransferase refers to any enzyme which catalyzes the conversion of homocysteine via methylation to methionine.

As used herein, adenosylhomocysteinase refers to any enzyme that catalyzes the ATP-dependent conversion of S-adenosylmethionine (AdoMet) to methylthioadenosine and L-homoserine.

As used herein, cystathionine  $\beta$ -synthase refers to any enzyme that catalyzes the conversion of homocysteine and serine to cystathionine.

As used herein, cystathionine  $\gamma$ -lyase refers to any enzyme that catalyzes the  $\gamma$  cleavage of cystathionine.

As used herein, *O*-acetylhomoserine (thiol)-lyase refers to any enzyme that catalyzes the conversion of *O*-acetylhomoserine and sulfur to homocysteine.

## **Agents**

### **(a) Nucleic Acid Molecules**

Agents of the present invention include plant nucleic acid molecules and more specifically include maize and soybean nucleic acid molecules and more specifically include nucleic acid molecules of the maize genotypes B73 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), B73 x Mo17 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), DK604 (Dekalb Genetics, Dekalb, Illinois U.S.A.), H99 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), RX601 (Asgrow Seed Company, Des Moines, Iowa), Mo17 (Illinois Foundation Seeds, Champaign, Illinois U.S.A.), and soybean types Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa), C1944 (United States Department of Agriculture (USDA) Soybean Germplasm Collection, Urbana, Illinois U.S.A.), Cristalina (USDA Soybean Germplasm Collection, Urbana,

Illinois U.S.A.), FT108 (Monsoy, Brazil), Hartwig (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.), BW211S Null (Tohoku University, Morioka, Japan), PI507354 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.), Asgrow A4922 (Asgrow Seed Company, Des Moines, Iowa U.S.A.), PI227687 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.), PI229358 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and Asgrow A3237 (Asgrow Seed Company, Des Moines, Iowa U.S.A.).

A subset of the nucleic acid molecules of the present invention includes nucleic acid molecules that are marker molecules. Another subset of the nucleic acid molecules of the present invention include nucleic acid molecules that encode a protein or fragment thereof. Another subset of the nucleic acid molecules of the present invention are EST molecules.

Fragment nucleic acid molecules may encode significant portion(s) of, or indeed most of, these nucleic acid molecules. Alternatively, the fragments may comprise smaller oligonucleotides (having from about 15 to about 250 nucleotide residues and more preferably, about 15 to about 30 nucleotide residues).

As used herein, an agent, be it a naturally occurring molecule or otherwise may be “substantially purified,” if desired, such that one or more molecules that is or may be present in a naturally occurring preparation containing that molecule will have been removed or will be present at a lower concentration than that at which it would normally be found.

The agents of the present invention will preferably be “biologically active” with respect to either a structural attribute, such as the capacity of a nucleic acid to hybridize to another nucleic acid molecule, or the ability of a protein to be bound by an antibody (or to compete with another molecule for such binding). Alternatively, such an attribute may be catalytic and thus involve the capacity of the agent to mediate a chemical reaction or response.

The agents of the present invention may also be recombinant. As used herein, the term recombinant means any agent (e.g. DNA, peptide etc.), that is, or results, however indirect, from human manipulation of a nucleic acid molecule.

It is understood that the agents of the present invention may be labeled with reagents that facilitate detection of the agent (e.g. fluorescent labels, Prober *et al.*, *Science* 238:336-340 (1987); Albarella *et al.*, EP 144914; chemical labels, Sheldon *et al.*, U.S. Patent 4,582,789; Albarella *et al.*, U.S. Patent 4,563,417; modified bases, Miyoshi *et al.*, EP 119448, all of which are hereby incorporated by reference in their entirety).

It is further understood, that the present invention provides recombinant bacterial, mammalian, microbial, insect, fungal and plant cells and viral constructs comprising the agents of the present invention. (See, for example, Uses of the Agents of the Invention, Section (a) Plant Constructs and Plant Transformants; Section (b) Fungal Constructs and Fungal Transformants; Section (c) Mammalian Constructs and Transformed Mammalian Cells; Section (d) Insect Constructs and Transformed Insect Cells; and Section (e) Bacterial Constructs and Transformed Bacterial Cells)

Nucleic acid molecules or fragments thereof of the present invention are capable of specifically hybridizing to other nucleic acid molecules under certain circumstances. As used herein, two nucleic acid molecules are said to be capable of specifically hybridizing to one another if the two molecules are capable of forming an anti-parallel, double-stranded nucleic acid structure. A nucleic acid molecule is said to be the “complement” of another nucleic acid molecule if they exhibit complete complementarity. As used herein, molecules are said to exhibit “complete complementarity” when every nucleotide of one of the molecules is complementary to a nucleotide of the other. Two molecules are said to be “minimally



complementary” if they can hybridize to one another with sufficient stability to permit them to remain annealed to one another under at least conventional "low-stringency" conditions.

Similarly, the molecules are said to be “complementary” if they can hybridize to one another with sufficient stability to permit them to remain annealed to one another under conventional "high-stringency" conditions. Conventional stringency conditions are described by Sambrook *et al.*, *Molecular Cloning*, A Laboratory Manual, 2nd Ed., Cold Spring Harbor Press, Cold Spring Harbor, New York (1989) and by Haymes *et al.*, *Nucleic Acid Hybridization, A Practical Approach*, IRL Press, Washington, DC (1985), the entirety of which is herein incorporated by reference. Departures from complete complementarity are therefore permissible, as long as such departures do not completely preclude the capacity of the molecules to form a double-stranded structure. Thus, in order for a nucleic acid molecule to serve as a primer or probe it need only be sufficiently complementary in sequence to be able to form a stable double-stranded structure under the particular solvent and salt concentrations employed.

Appropriate stringency conditions which promote DNA hybridization, for example, 6.0 X sodium chloride/sodium citrate (SSC) at about 45°C, followed by a wash of 2.0 X SSC at 50°C, are known to those skilled in the art or can be found in *Current Protocols in Molecular Biology*, John Wiley & Sons, N.Y. (1989), 6.3.1-6.3.6. For example, the salt concentration in the wash step can be selected from a low stringency of about 2.0 X SSC at 50°C to a high stringency of about 0.2 X SSC at 50°C. In addition, the temperature in the wash step can be increased from low stringency conditions at room temperature, about 22°C, to high stringency conditions at about 65°C. Both temperature and salt may be varied, or either the temperature or the salt concentration may be held constant while the other variable is changed.

In a preferred embodiment, a nucleic acid of the present invention will specifically hybridize to one or more of the nucleic acid molecules set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof under moderately stringent conditions, for example at about 2.0 X SSC and about 65°C.

In a particularly preferred embodiment, a nucleic acid of the present invention will include those nucleic acid molecules that specifically hybridize to one or more of the nucleic acid molecules set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof under high stringency conditions such as 0.2 X SSC and about 65°C.

In one aspect of the present invention, the nucleic acid molecules of the present invention have one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In another aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 90% sequence identity with one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In a further aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 95% sequence identity with one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In a more preferred aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 98% sequence identity with one or more of the nucleic acid sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof. In an even more preferred aspect of the present invention, one or more of the nucleic acid molecules of the present invention share between 100% and 99% sequence identity with one or more of the sequences set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof.

In a further more preferred aspect of the present invention, one or more of the nucleic acid molecules of the present invention exhibit 100% sequence identity with a nucleic acid molecule present within MONN01, SATMON001 through SATMON031, SATMON033, SATMON034, SATMON~001, SATMONN01, SATMONN04 through SATMONN006, CMz029 through CMz031, CMz033, CMz035 through CMz037, CMz039 through CMz042, CMz044 through CMz045, CMz047 through CMz050, SOYMON001 through SOYMON038, Soy51 through Soy56, Soy58 through Soy62, Soy65 through Soy66, Soy 68 through Soy73 and Soy76 through Soy77, Lib9, Lib22 through Lib25, Lib35, Lib80 through Lib81, Lib 144, Lib146, Lib147, Lib190, Lib3032 through Lib3036 and Lib3099 (Monsanto Company, St. Louis, Missouri U.S.A.).

**(i) Nucleic Acid Molecules Encoding Proteins or Fragments Thereof**

Nucleic acid molecules of the present invention can comprise sequences that encode a methionine pathway protein or fragment thereof. Such proteins or fragments thereof include homologues of known proteins in other organisms.

In a preferred embodiment of the present invention, a maize or soybean protein homologue or fragment thereof of the present invention is a homologue of another plant protein. In another preferred embodiment of the present invention, a maize or soybean protein homologue or fragment thereof of the present invention is a homologue of a fungal protein. In another preferred embodiment of the present invention, a maize or soybean protein homologue of the present invention is a homologue of mammalian protein. In another preferred embodiment of the present invention, a maize or soybean protein homologue or fragment thereof of the present invention is a homologue of a bacterial protein. In another preferred embodiment of the present invention, a soybean protein homologue or fragment thereof of the present invention is a

maize protein homologue or fragment thereof of the present invention is a homologue of a soybean protein.

In a preferred embodiment of the present invention, the nucleic molecule of the present invention encodes a maize or soybean homologue protein or fragment thereof where a maize or soybean homologue protein exhibits a BLAST probability score of greater than 1E-12, preferably a BLAST probability score of between about 1E-30 and about 1E-12, even more preferably a BLAST probability score of greater than 1E-30 with its homologue.

In another preferred embodiment of the present invention, the nucleic acid molecule encoding a maize or soybean protein homologue or fragment thereof or fragment thereof exhibits a % identity with its homologue of between about 25% and about 40%, more preferably of between about 40 and about 70%, even more preferably of between about 70% and about 90% and even more preferably between about 90% and 99%. In another preferred embodiment, of the present invention, a maize or soybean protein homologue or fragments thereof exhibits a % identity with its homologue of 100%.

In a preferred embodiment of the present invention, the nucleic molecule of the present invention encodes a maize or soybean homologue protein or fragment thereof where a maize or soybean homologue protein exhibits a BLAST score of greater than 120, preferably a BLAST score of between about 1450 and about 120, even more preferably a BLAST score of greater than 1450 with its homologue.

Nucleic acid molecules of the present invention also include non-maize, non-soybean homologues. Preferred non- homologues are selected from the group consisting of alfalfa, *Arabidopsis*, barley, *Brassica*, broccoli, cabbage, citrus, cotton, garlic, oat, oilseed rape, onion, canola, flax, an ornamental plant, pea, peanut, pepper, potato, rice, rye, sorghum, strawberry,

sugarcane, sugarbeet, tomato, wheat, poplar, pine, fir, eucalyptus, apple, lettuce, lentils, grape, banana, tea, turf grasses, sunflower, oil palm and *Phaseolus*.

In a preferred embodiment, nucleic acid molecules having SEQ ID NO: 1 through SEQ ID NO: 3204 or complements and fragments of either can be utilized to obtain such homologues.

The degeneracy of the genetic code, which allows different nucleic acid sequences to code for the same protein or peptide, is known in the literature. (U.S. Patent No. 4,757,006, the entirety of which is herein incorporated by reference).

In an aspect of the present invention, one or more of the nucleic acid molecules of the present invention differ in nucleic acid sequence from those encoding maize or soybean homologue or fragment thereof in SEQ ID NO: 1 through SEQ ID NO: 3204 due to the degeneracy in the genetic code in that they encode the same protein but differ in nucleic acid sequence.

In another further aspect of the present invention, one or more of the nucleic acid molecules of the present invention differ in nucleic acid sequence from those encoding maize or soybean homologue or fragment thereof in SEQ ID NO: 1 through SEQ ID NO: 3204 due to fact that the different nucleic acid sequence encodes a protein having one or more conservative amino acid residue. Examples of conservative substitutions are set forth in Table 1. It is understood that codons capable of coding for such conservative substitutions are known in the art.

**Table 1**

<u>Original Residue</u>	<u>Conservative Substitutions</u>
Ala	Ser
Arg	Lys

Asn	Gln; His
Asp	Glu
Cys	Ser; Ala
Gln	Asn
Glu	Asp
Gly	Pro
His	Asn; Gln
Ile	Leu; Val
Leu	Ile; Val
Lys	Arg; Gln; Glu
Met	Leu; Ile
Phe	Met; Leu; Tyr
Ser	Thr
Thr	Ser
Trp	Tyr
Tyr	Trp; Phe
Val	Ile; Leu

In a further aspect of the present invention, one or more of the nucleic acid molecules of the present invention differ in nucleic acid sequence from those encoding a maize or soybean homologue or fragment thereof set forth in SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof due to the fact that one or more codons encoding an amino acid has been

substituted for a codon that encodes a nonessential substitution of the amino acid originally encoded.

Agents of the present invention include nucleic acid molecules that encode a maize or soybean methionine pathway protein or fragment thereof and particularly substantially purified nucleic acid molecules selected from the group consisting of a nucleic acid molecule that encodes a maize or soybean methionine adenosyltransferase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean S-adenosylmethionine decarboxylase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean aspartate kinase protein or fragment thereof, a nucleic acid molecule that encode a maize or soybean aspartate-semialdehyde dehydrogenase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean *O*-succinylhomoserine (thiol)-lyase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean cystathionine  $\beta$ -lyase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean adenosylhomocysteine protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean cystathionine  $\beta$ -synthase protein or fragment thereof, a nucleic acid molecule that encodes a maize or soybean cystathionine  $\gamma$ -lyase protein or fragment thereof, and a nucleic acid molecule that encodes a maize or soybean *O*-acetylhomoserine (thiol)-lyase protein or fragment thereof.

Non-limiting examples of such nucleic acid molecules of the present invention are nucleic acid molecules comprising: SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof that encode for a methionine pathway protein or fragment thereof, SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479 or fragment thereof that

encode for a methionine adenosyltransferase protein or fragment thereof, SEQ ID NO: 430  
 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623 or fragment thereof  
 that encode for a S-adenosylmethionine decarboxylase protein or fragment thereof, SEQ ID NO:  
 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648 or fragment  
 thereof that encode for a aspartate kinase protein or fragment thereof, SEQ ID NO: 901 through  
 SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654 or fragment thereof that  
 encode for a aspartate-semialdehyde dehydrogenase protein or fragment thereof, SEQ ID NO:  
 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or fragment  
 thereof that encode for a *O*-succinylhomoserine (thiol)-lyase protein or fragment thereof, SEQ ID  
 NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or  
 fragment thereof that encode for a cystathionine  $\beta$ -lyase protein or fragment thereof, SEQ ID  
 NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992 or  
 fragment thereof that encode for a 5-methyltetrahydropteroyltriglutamate-homocysteine-S-  
 methyltransferase protein or fragment thereof, SEQ ID NO: 1354 through SEQ ID NO: 1630 and  
 SEQ ID NO: 2993 through SEQ ID NO: 3199 or fragment thereof that encode for an  
 adenosylhomocysteinase protein or fragment thereof, SEQ ID NO: 1631 through SEQ ID NO:  
 1632 or fragment thereof that encode for a cystathionine  $\beta$ -synthase protein or fragment thereof,  
 SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204  
 or fragment thereof that encode for a cystathionine  $\gamma$ -lyase protein or fragment thereof, and SEQ  
 ID NO: 3200 through SEQ ID NO: 3202 or fragment thereof that encode for an *O*-  
 acetylhomoserine (thiol)-lyase protein or fragment thereof.



A nucleic acid molecule of the present invention can also encode an homologue of a maize or soybean methionine adenosyltransferase or fragment thereof, a maize or soybean S-adenosylmethionine decarboxylase or fragment thereof, a maize or soybean aspartate kinase or fragment thereof, a maize or soybean aspartate-semialdehyde dehydrogenase or fragment thereof, a maize or soybean *O*-succinylhomoserine (thiol)-lyase or fragment thereof, a maize or soybean cystathionine  $\beta$ -lyase or fragment thereof, a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof, a maize or soybean adenosylhomocysteinease or fragment thereof, a maize or soybean cystathionine  $\beta$ -synthase or fragment thereof, a maize or soybean cystathionine  $\gamma$ -lyase or fragment thereof or a maize or soybean *O*-acetylhomoserine (thiol)-lyase or fragment thereof. As used herein a homologue protein molecule or fragment thereof is a counterpart protein molecule or fragment thereof in a second species (*e.g.*, maize methionine adenosyltransferase protein is a homologue of *Arabidopsis*' methionine adenosyltransferase protein).

## **(ii) Nucleic Acid Molecule Markers and Probes**

One aspect of the present invention concerns markers that include nucleic acid molecules SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragments of either that can act as markers. Genetic markers of the present invention include “dominant” or “codominant” markers. “Codominant markers” reveal the presence of two or more alleles (two per diploid individual) at a locus. “Dominant markers” reveal the presence of only a single allele per locus. The presence of the dominant marker phenotype (*e.g.*, a band of DNA) is an indication that one allele is present in either the homozygous or heterozygous condition. The absence of the dominant marker phenotype (*e.g.* absence of a DNA band) is merely evidence that “some other”

undefined allele is present. In the case of populations where individuals are predominantly homozygous and loci are predominately dimorphic, dominant and codominant markers can be equally valuable. As populations become more heterozygous and multi-allelic, codominant markers often become more informative of the genotype than dominant markers. Marker molecules can be, for example, capable of detecting polymorphisms such as single nucleotide polymorphisms (SNPs).

SNPs are single base changes in genomic DNA sequence. They occur at greater frequency and are spaced with a greater uniformity throughout a genome than other reported forms of polymorphism. The greater frequency and uniformity of SNPs means that there is greater probability that such a polymorphism will be found near or in a genetic locus of interest than would be the case for other polymorphisms. SNPs are located in protein-coding regions and noncoding regions of a genome. Some of these SNPs may result in defective or variant protein expression (e.g., as a results of mutations or defective splicing). Analysis (genotyping) of characterized SNPs can require only a plus/minus assay rather than a lengthy measurement, permitting easier automation.

SNPs can be characterized using any of a variety of methods. Such methods include the direct or indirect sequencing of the site, the use of restriction enzymes (Botstein *et al.*, *Am. J. Hum. Genet.* 32:314-331 (1980), the entirety of which is herein incorporated reference; Konieczny and Ausubel, *Plant J.* 4:403-410 (1993), the entirety of which is herein incorporated by reference), enzymatic and chemical mismatch assays (Myers *et al.*, *Nature* 313:495-498 (1985), the entirety of which is herein incorporated by reference), allele-specific PCR (Newton *et al.*, *Nucl. Acids Res.* 17:2503-2516 (1989), the entirety of which is herein incorporated by reference; Wu *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:2757-2760 (1989), the entirety of which

is herein incorporated by reference), ligase chain reaction (Barany, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:189-193 (1991), the entirety of which is herein incorporated by reference), single-strand conformation polymorphism analysis (Labrune *et al.*, *Am. J. Hum. Genet.* 48: 1115-1120 (1991), the entirety of which is herein incorporated by reference), primer-directed nucleotide incorporation assays (Kuppuswami *et al.*, *Proc. Natl. Acad. Sci. USA* 88:1143-1147 (1991), the entirety of which is herein incorporated by reference), dideoxy fingerprinting (Sarkar *et al.*, *Genomics* 13:441-443 (1992), the entirety of which is herein incorporated by reference), solid-phase ELISA-based oligonucleotide ligation assays (Nikiforov *et al.*, *Nucl. Acids Res.* 22:4167-4175 (1994), the entirety of which is herein incorporated by reference), oligonucleotide fluorescence-quenching assays (Livak *et al.*, *PCR Methods Appl.* 4:357-362 (1995), the entirety of which is herein incorporated by reference), 5'-nuclease allele-specific hybridization TaqMan assay (Livak *et al.*, *Nature Genet.* 9:341-342 (1995), the entirety of which is herein incorporated by reference), template-directed dye-terminator incorporation (TDI) assay (Chen and Kwok, *Nucl. Acids Res.* 25:347-353 (1997), the entirety of which is herein incorporated by reference), allele-specific molecular beacon assay (Tyagi *et al.*, *Nature Biotech.* 16: 49-53 (1998), the entirety of which is herein incorporated by reference), PinPoint assay (Haff and Smirnov, *Genome Res.* 7: 378-388 (1997), the entirety of which is herein incorporated by reference) and dCAPS analysis (Neff *et al.*, *Plant J.* 14:387-392 (1998), the entirety of which is herein incorporated by reference).

Additional markers, such as AFLP markers, RFLP markers and RAPD markers, can be utilized (Walton, *Seed World* 22-29 (July, 1993), the entirety of which is herein incorporated by reference; Burow and Blake, *Molecular Dissection of Complex Traits*, 13-29, Paterson (ed.), CRC Press, New York (1988), the entirety of which is herein incorporated by reference). DNA

markers can be developed from nucleic acid molecules using restriction endonucleases, the PCR and/or DNA sequence information. RFLP markers result from single base changes or insertions/deletions. These codominant markers are highly abundant in plant genomes, have a medium level of polymorphism and are developed by a combination of restriction endonuclease digestion and Southern blotting hybridization. CAPS are similarly developed from restriction nuclease digestion but only of specific PCR products. These markers are also codominant, have a medium level of polymorphism and are highly abundant in the genome. The CAPS result from single base changes and insertions/deletions.

Another marker type, RAPDs, are developed from DNA amplification with random primers and result from single base changes and insertions/deletions in plant genomes. They are dominant markers with a medium level of polymorphisms and are highly abundant. AFLP markers require using the PCR on a subset of restriction fragments from extended adapter primers. These markers are both dominant and codominant are highly abundant in genomes and exhibit a medium level of polymorphism.

SSRs require DNA sequence information. These codominant markers result from repeat length changes, are highly polymorphic and do not exhibit as high a degree of abundance in the genome as CAPS, AFLPs and RAPDs SNPs also require DNA sequence information. These codominant markers result from single base substitutions. They are highly abundant and exhibit a medium of polymorphism (Rafalski *et al.*, In: *Nonmammalian Genomic Analysis*, Birren and Lai (ed.), Academic Press, San Diego, CA, pp. 75-134 (1996), the entirety of which is herein incorporated by reference). It is understood that a nucleic acid molecule of the present invention may be used as a marker.

A PCR probe is a nucleic acid molecule capable of initiating a polymerase activity while in a double-stranded structure to with another nucleic acid. Various methods for determining the structure of PCR probes and PCR techniques exist in the art. Computer generated searches using programs such as Primer3 ([www-genome.wi.mit.edu/cgi-bin/primer/primer3.cgi](http://www-genome.wi.mit.edu/cgi-bin/primer/primer3.cgi)), STSPipeline ([www-genome.wi.mit.edu/cgi-bin/www-STS\\_Pipeline](http://www-genome.wi.mit.edu/cgi-bin/www-STS_Pipeline)), or GeneUp (Pesole *et al.*, *BioTechniques* 25:112-123 (1998) the entirety of which is herein incorporated by reference), for example, can be used to identify potential PCR primers.

It is understood that a fragment of one or more of the nucleic acid molecules of the present invention may be a probe and specifically a PCR probe.

#### **(b) Protein and Peptide Molecules**

A class of agents comprises one or more of the protein or fragments thereof or peptide molecules encoded by SEQ ID NO: 1 through SEQ ID NO: 3204 or one or more of the protein or fragment thereof and peptide molecules encoded by other nucleic acid agents of the present invention. As used herein, the term "protein molecule" or "peptide molecule" includes any molecule that comprises five or more amino acids. It is well known in the art that proteins may undergo modification, including post-translational modifications, such as, but not limited to, disulfide bond formation, glycosylation, phosphorylation, or oligomerization. Thus, as used herein, the term "protein molecule" or "peptide molecule" includes any protein molecule that is modified by any biological or non-biological process. The terms "amino acid" and "amino acids" refer to all naturally occurring L-amino acids. This definition is meant to include norleucine, ornithine, homocysteine and homoserine.

Non-limiting examples of the protein or fragment thereof of the present invention include a maize or soybean methionine pathway protein or fragment thereof; a maize or soybean

methionine adenosyltransferase or fragment thereof, a maize or soybean S-adenosylmethionine decarboxylase or fragment thereof, a maize or soybean aspartate kinase or fragment thereof, a maize or soybean aspartate-semialdehyde dehydrogenase or fragment thereof, a maize or soybean *O*-succinylhomoserine (thiol)-lyase or fragment thereof, a maize or soybean cystathionine  $\beta$ -lyase or fragment thereof, a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof, a maize or soybean adenosylhomocysteinase or fragment thereof, a maize or soybean cystathionine  $\beta$ -synthase or fragment thereof, a maize or soybean cystathionine  $\gamma$ -lyase or fragment thereof or a maize or soybean *O*-acetylhomoserine (thiol)-lyase or fragment thereof.

Non-limiting examples of the protein or fragment molecules of the present invention are an methionine pathway protein or fragment thereof encoded by: SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof that encode for a methionine pathway protein or fragment thereof, SEQ ID NO: 1 through SEQ ID NO: 429 and SEQ ID NO: 1635 through SEQ ID NO: 2479 or fragment thereof that encode for a methionine adenosyltransferase protein or fragment thereof, SEQ ID NO: 430 through SEQ ID NO: 857 and SEQ ID NO: 2480 through SEQ ID NO: 2623 or fragment thereof that encode for a S-adenosylmethionine decarboxylase protein or fragment thereof, SEQ ID NO: 858 through SEQ ID NO: 900 and SEQ ID NO: 2624 through SEQ ID NO: 2648 or fragment thereof that encode for a aspartate kinase protein or fragment thereof, SEQ ID NO: 901 through SEQ ID NO: 904 and SEQ ID NO: 2649 through SEQ ID NO: 2654 or fragment thereof that encode for a aspartate-semialdehyde dehydrogenase protein or fragment thereof, SEQ ID NO: 905 through SEQ ID NO: 953 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or fragment thereof that encode for a *O*-succinylhomoserine (thiol)-lyase protein or

fragment thereof, SEQ ID NO: 954 through SEQ ID NO: 963 and SEQ ID NO: 2655 through SEQ ID NO: 2660 or fragment thereof that encode for a cystathionine  $\beta$ -lyase or fragment thereof, SEQ ID NO: 964 through SEQ ID NO: 1353 and SEQ ID NO: 2666 through SEQ ID NO: 2992 or fragment thereof that encode for a 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase protein or fragment thereof, SEQ ID NO: 1354 through SEQ ID NO: 1630 and SEQ ID NO: 2993 through SEQ ID NO: 3199 or fragment thereof that encode for an adenosylhomocysteinase protein or fragment thereof, SEQ ID NO: 1631 through SEQ ID NO: 1632 or fragment thereof that encode for a cystathionine  $\beta$ -synthase protein or fragment thereof, SEQ ID NO: 1633 through SEQ ID NO: 1634 and SEQ ID NO: 3203 through SEQ ID NO: 3204 or fragment thereof that encode for a cystathionine  $\gamma$ -lyase protein or fragment thereof, and SEQ ID NO: 3200 through SEQ ID NO: 3202 or fragment thereof that encode for an *O*-acetylhomoserine (thiol)-lyase protein or fragment thereof.

One or more of the protein or fragment of peptide molecules may be produced via chemical synthesis, or more preferably, by expressing in a suitable bacterial or eucaryotic host. Suitable methods for expression are described by Sambrook *et al.*, (In: *Molecular Cloning, A Laboratory Manual, 2nd Edition, Cold Spring Harbor Press, Cold Spring Harbor, New York* (1989)), or similar texts. For example, the protein may be expressed in, for example, Uses of the Agents of the Invention, Section (a) Plant Constructs and Plant Transformants; Section (b) Fungal Constructs and Fungal Transformants; Section (c) Mammalian Constructs and Transformed Mammalian Cells; Section (d) Insect Constructs and Transformed Insect Cells; and Section (e) Bacterial Constructs and Transformed Bacterial Cells.

A “protein fragment” is a peptide or polypeptide molecule whose amino acid sequence comprises a subset of the amino acid sequence of that protein. A protein or fragment thereof that comprises one or more additional peptide regions not derived from that protein is a “fusion” protein. Such molecules may be derivatized to contain carbohydrate or other moieties (such as keyhole limpet hemocyanin, etc.). Fusion protein or peptide molecules of the present invention are preferably produced via recombinant means.

Another class of agents comprise protein or peptide molecules or fragments or fusions thereof encoded by SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof in which conservative, non-essential or non-relevant amino acid residues have been added, replaced or deleted. Computerized means for designing modifications in protein structure are known in the art (Dahiyat and Mayo, *Science* 278:82-87 (1997), the entirety of which is herein incorporated by reference).

The protein molecules of the present invention include plant homologue proteins. An example of such a homologue is a homologue protein of a non-maize or non soybean plant species, that include but not limited to alfalfa, *Arabidopsis*, barley, *Brassica*, broccoli, cabbage, citrus, cotton, garlic, oat, oilseed rape, onion, canola, flax, an ornamental plant, pea, peanut, pepper, potato, rice, rye, sorghum, strawberry, sugarcane, sugarbeet, tomato, wheat, poplar, pine, fir, eucalyptus, apple, lettuce, lentils, grape, banana, tea, turf grasses, sunflower, oil palm, *Phaseolus* etc. Particularly preferred non-maize or non-soybean for use for the isolation of homologs would include, *Arabidopsis*, barley, cotton, oat, oilseed rape, rice, canola, ornamentals, sugarcane, sugarbeet, tomato, potato, wheat and turf grasses. Such a homologue can be obtained by any of a variety of methods. Most preferably, as indicated above, one or more of the disclosed sequences (SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof) will be



used to define a pair of primers that may be used to isolate the homologue-encoding nucleic acid molecules from any desired species. Such molecules can be expressed to yield homologues by recombinant means.

### **(c) Antibodies**

One aspect of the present invention concerns antibodies, single-chain antigen binding molecules, or other proteins that specifically bind to one or more of the protein or peptide molecules of the present invention and their homologues, fusions or fragments. Such antibodies may be used to quantitatively or qualitatively detect the protein or peptide molecules of the present invention. As used herein, an antibody or peptide is said to “specifically bind” to a protein or peptide molecule of the present invention if such binding is not competitively inhibited by the presence of non-related molecules.

Nucleic acid molecules that encode all or part of the protein of the present invention can be expressed, via recombinant means, to yield protein or peptides that can in turn be used to elicit antibodies that are capable of binding the expressed protein or peptide. Such antibodies may be used in immunoassays for that protein. Such protein-encoding molecules, or their fragments may be a “fusion” molecule (i.e., a part of a larger nucleic acid molecule) such that, upon expression, a fusion protein is produced. It is understood that any of the nucleic acid molecules of the present invention may be expressed, via recombinant means, to yield proteins or peptides encoded by these nucleic acid molecules.

The antibodies that specifically bind proteins and protein fragments of the present invention may be polyclonal or monoclonal and may comprise intact immunoglobulins, or antigen binding portions of immunoglobulins fragments (such as  $F(ab')$ ,  $F(ab')_2$ ), or single-chain

immunoglobulins producible, for example, via recombinant means. It is understood that practitioners are familiar with the standard resource materials which describe specific conditions and procedures for the construction, manipulation and isolation of antibodies (*see*, for example, Harlow and Lane, In: *Antibodies: A Laboratory Manual*, Cold Spring Harbor Press, Cold Spring Harbor, New York (1988), the entirety of which is herein incorporated by reference).

Murine monoclonal antibodies are particularly preferred. BALB/c mice are preferred for this purpose, however, equivalent strains may also be used. The animals are preferably immunized with approximately 25 µg of purified protein (or fragment thereof) that has been emulsified in a suitable adjuvant (such as TiterMax adjuvant (Vaxcel, Norcross, GA)). Immunization is preferably conducted at two intramuscular sites, one intraperitoneal site and one subcutaneous site at the base of the tail. An additional i.v. injection of approximately 25 µg of antigen is preferably given in normal saline three weeks later. After approximately 11 days following the second injection, the mice may be bled and the blood screened for the presence of anti-protein or peptide antibodies. Preferably, a direct binding Enzyme-Linked Immunoassay (ELISA) is employed for this purpose.

More preferably, the mouse having the highest antibody titer is given a third i.v. injection of approximately 25 µg of the same protein or fragment. The splenic leukocytes from this animal may be recovered 3 days later and then permitted to fuse, most preferably, using polyethylene glycol, with cells of a suitable myeloma cell line (such as, for example, the P3X63Ag8.653 myeloma cell line). Hybridoma cells are selected by culturing the cells under “HAT” (hypoxanthine-aminopterin-thymine) selection for about one week. The resulting clones may then be screened for their capacity to produce monoclonal antibodies (“mAbs”), preferably by direct ELISA.

In one embodiment, anti-protein or peptide monoclonal antibodies are isolated using a fusion of a protein or peptide of the present invention, or conjugate of a protein or peptide of the present invention, as immunogens. Thus, for example, a group of mice can be immunized using a fusion protein emulsified in Freund's complete adjuvant (*e.g.* approximately 50 µg of antigen per immunization). At three week intervals, an identical amount of antigen is emulsified in Freund's incomplete adjuvant and used to immunize the animals. Ten days following the third immunization, serum samples are taken and evaluated for the presence of antibody. If antibody titers are too low, a fourth booster can be employed. Polysera capable of binding the protein or peptide can also be obtained using this method.

In a preferred procedure for obtaining monoclonal antibodies, the spleens of the above-described immunized mice are removed, disrupted and immune splenocytes are isolated over a ficoll gradient. The isolated splenocytes are fused, using polyethylene glycol with BALB/c-derived HGPRT (hypoxanthine guanine phosphoribosyl transferase) deficient P3x63xAg8.653 plasmacytoma cells. The fused cells are plated into 96 well microtiter plates and screened for hybridoma fusion cells by their capacity to grow in culture medium supplemented with hypoxanthine, aminopterin and thymidine for approximately 2-3 weeks.

Hybridoma cells that arise from such incubation are preferably screened for their capacity to produce an immunoglobulin that binds to a protein of interest. An indirect ELISA may be used for this purpose. In brief, the supernatants of hybridomas are incubated in microtiter wells that contain immobilized protein. After washing, the titer of bound immunoglobulin can be determined using, for example, a goat anti-mouse antibody conjugated to horseradish peroxidase. After additional washing, the amount of immobilized enzyme is determined (for example through the use of a chromogenic substrate). Such screening is performed as quickly as possible

after the identification of the hybridoma in order to ensure that a desired clone is not overgrown by non-secreting neighbor cells. Desirably, the fusion plates are screened several times since the rates of hybridoma growth vary. In a preferred sub-embodiment, a different antigenic form may be used to screen the hybridoma. Thus, for example, the splenocytes may be immunized with one immunogen, but the resulting hybridomas can be screened using a different immunogen. It is understood that any of the protein or peptide molecules of the present invention may be used to raise antibodies.

As discussed below, such antibody molecules or their fragments may be used for diagnostic purposes. Where the antibodies are intended for diagnostic purposes, it may be desirable to derivatize them, for example with a ligand group (such as biotin) or a detectable marker group (such as a fluorescent group, a radioisotope or an enzyme).

The ability to produce antibodies that bind the protein or peptide molecules of the present invention permits the identification of mimetic compounds of those molecules. A “mimetic compound” is a compound that is not that compound, or a fragment of that compound, but which nonetheless exhibits an ability to specifically bind to antibodies directed against that compound.

It is understood that any of the agents of the present invention can be substantially purified and/or be biologically active and/or recombinant.

### **Uses of the Agents of the Invention**

Nucleic acid molecules and fragments thereof of the present invention may be employed to obtain other nucleic acid molecules from the same species (e.g., ESTs or fragment thereof from maize may be utilized to obtain other nucleic acid molecules from maize). Such nucleic acid molecules include the nucleic acid molecules that encode the complete coding sequence of a protein and promoters and flanking sequences of such molecules. In addition, such nucleic acid

molecules include nucleic acid molecules that encode for other isozymes or gene family members. Such molecules can be readily obtained by using the above-described nucleic acid molecules or fragments thereof to screen cDNA or genomic libraries obtained from maize or soybean. Methods for forming such libraries are well known in the art.

Nucleic acid molecules and fragments thereof of the present invention may also be employed to obtain nucleic acid homologues. Such homologues include the nucleic acid molecule of other plants or other organisms (*e.g.*, alfalfa, *Arabidopsis*, barley, *Brassica*, broccoli, cabbage, citrus, cotton, garlic, oat, oilseed rape, onion, canola, flax, an ornamental plant, pea, peanut, pepper, potato, rice, rye, sorghum, strawberry, sugarcane, sugarbeet, tomato, wheat, poplar, pine, fir, eucalyptus, apple, lettuce, lentils, grape, banana, tea, turf grasses, sunflower, oil palm, *Phaseolus*, etc.) including the nucleic acid molecules that encode, in whole or in part, protein homologues of other plant species or other organisms, sequences of genetic elements such as promoters and transcriptional regulatory elements. Such molecules can be readily obtained by using the above-described nucleic acid molecules or fragments thereof to screen cDNA or genomic libraries obtained from such plant species. Methods for forming such libraries are well known in the art. Such homologue molecules may differ in their nucleotide sequences from those found in one or more of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof because complete complementarity is not needed for stable hybridization. The nucleic acid molecules of the present invention therefore also include molecules that, although capable of specifically hybridizing with the nucleic acid molecules may lack “complete complementarity.”

Any of a variety of methods may be used to obtain one or more of the above-described nucleic acid molecules (Zamechik *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 83:4143-4146 (1986), the entirety of which is herein incorporated by reference; Goodchild *et al.*, *Proc. Natl. Acad. Sci.*

(U.S.A.) 85:5507-5511 (1988), the entirety of which is herein incorporated by reference; Wickstrom *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 85:1028-1032 (1988), the entirety of which is herein incorporated by reference; Holt *et al.*, *Molec. Cell. Biol.* 8:963-973 (1988), the entirety of which is herein incorporated by reference; Gerwartz *et al.*, *Science* 242:1303-1306 (1988), the entirety of which is herein incorporated by reference; Anfossi *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:3379-3383 (1989), the entirety of which is herein incorporated by reference; Becker *et al.*, *EMBO J.* 8:3685-3691 (1989); the entirety of which is herein incorporated by reference).

Automated nucleic acid synthesizers may be employed for this purpose. In lieu of such synthesis, the disclosed nucleic acid molecules may be used to define a pair of primers that can be used with the polymerase chain reaction (Mullis *et al.*, *Cold Spring Harbor Symp. Quant. Biol.* 51:263-273 (1986); Erlich *et al.*, European Patent 50,424; European Patent 84,796; European Patent 258,017; European Patent 237,362; Mullis, European Patent 201,184; Mullis *et al.*, U.S. Patent 4,683,202; Erlich, U.S. Patent 4,582,788; and Saiki *et al.*, U.S. Patent 4,683,194, all of which are herein incorporated by reference in their entirety) to amplify and obtain any desired nucleic acid molecule or fragment.

Promoter sequence(s) and other genetic elements, including but not limited to transcriptional regulatory flanking sequences, associated with one or more of the disclosed nucleic acid sequences can also be obtained using the disclosed nucleic acid sequence provided herein. In one embodiment, such sequences are obtained by incubating EST nucleic acid molecules or preferably fragments thereof with members of genomic libraries (*e.g.* maize and soybean) and recovering clones that hybridize to the EST nucleic acid molecule or fragment thereof. In a second embodiment, methods of "chromosome walking," or inverse PCR may be used to obtain such sequences (Frohman *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 85:8998-9002

(1988); Ohara *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:5673-5677 (1989); Pang *et al.*, *Biotechniques* 22:1046-1048 (1977); Huang *et al.*, *Methods Mol. Biol.* 69:89-96 (1997); Huang *et al.*, *Method Mol. Biol.* 67:287-294 (1997); Benkel *et al.*, *Genet. Anal.* 13:123-127 (1996); Hartl *et al.*, *Methods Mol. Biol.* 58:293-301 (1996), all of which are herein incorporated by reference in their entirety).

The nucleic acid molecules of the present invention may be used to isolate promoters of cell enhanced, cell specific, tissue enhanced, tissue specific, developmentally or environmentally regulated expression profiles. Isolation and functional analysis of the 5' flanking promoter sequences of these genes from genomic libraries, for example, using genomic screening methods and PCR techniques would result in the isolation of useful promoters and transcriptional regulatory elements. These methods are known to those of skill in the art and have been described (See, for example, Birren *et al.*, *Genome Analysis: Analyzing DNA*, 1, (1997), Cold Spring Harbor Laboratory Press, Cold Spring Harbor, N.Y., the entirety of which is herein incorporated by reference). Promoters obtained utilizing the nucleic acid molecules of the present invention could also be modified to affect their control characteristics. Examples of such modifications would include but are not limited to enhanced sequences as reported in Uses of the Agents of the Invention, Section (a) Plant Constructs and Plant Transformants. Such genetic elements could be used to enhance gene expression of new and existing traits for crop improvements.

In one sub-aspect, such an analysis is conducted by determining the presence and/or identity of polymorphism(s) by one or more of the nucleic acid molecules of the present invention and more preferably one or more of the EST nucleic acid molecule or fragment thereof which are associated with a phenotype, or a predisposition to that phenotype.

Any of a variety of molecules can be used to identify such polymorphism(s). In one embodiment, one or more of the EST nucleic acid molecules (or a sub-fragment thereof) may be employed as a marker nucleic acid molecule to identify such polymorphism(s). Alternatively, such polymorphisms can be detected through the use of a marker nucleic acid molecule or a marker protein that is genetically linked to (i.e., a polynucleotide that co-segregates with) such polymorphism(s).

In an alternative embodiment, such polymorphisms can be detected through the use of a marker nucleic acid molecule that is physically linked to such polymorphism(s). For this purpose, marker nucleic acid molecules comprising a nucleotide sequence of a polynucleotide located within 1mb of the polymorphism(s) and more preferably within 100kb of the polymorphism(s) and most preferably within 10kb of the polymorphism(s) can be employed.

The genomes of animals and plants naturally undergo spontaneous mutation in the course of their continuing evolution (Gusella, *Ann. Rev. Biochem.* 55:831-854 (1986)). A “polymorphism” is a variation or difference in the sequence of the gene or its flanking regions that arises in some of the members of a species. The variant sequence and the “original” sequence co-exist in the species’ population. In some instances, such co-existence is in stable or quasi-stable equilibrium.

A polymorphism is thus said to be “allelic,” in that, due to the existence of the polymorphism, some members of a species may have the original sequence (i.e., the original “allele”) whereas other members may have the variant sequence (i.e., the variant “allele”). In the simplest case, only one variant sequence may exist and the polymorphism is thus said to be di-allelic. In other cases, the species’ population may contain multiple alleles and the polymorphism is termed tri-allelic, etc. A single gene may have multiple different unrelated



polymorphisms. For example, it may have a di-allelic polymorphism at one site and a multi-allelic polymorphism at another site.

The variation that defines the polymorphism may range from a single nucleotide variation to the insertion or deletion of extended regions within a gene. In some cases, the DNA sequence variations are in regions of the genome that are characterized by short tandem repeats (STRs) that include tandem di- or tri-nucleotide repeated motifs of nucleotides. Polymorphisms characterized by such tandem repeats are referred to as "variable number tandem repeat" ("VNTR") polymorphisms. VNTRs have been used in identity analysis (Weber, U.S. Patent 5,075,217; Armour *et al.*, *FEBS Lett.* 307:113-115 (1992); Jones *et al.*, *Eur. J. Haematol.* 39:144-147 (1987); Horn *et al.*, PCT Patent Application WO91/14003; Jeffreys, European Patent Application 370,719; Jeffreys, U.S. Patent 5,175,082; Jeffreys *et al.*, *Amer. J. Hum. Genet.* 39:11-24 (1986); Jeffreys *et al.*, *Nature* 316:76-79 (1985); Gray *et al.*, *Proc. R. Acad. Soc. Lond.* 243:241-253 (1991); Moore *et al.*, *Genomics* 10:654-660 (1991); Jeffreys *et al.*, *Anim. Genet.* 18:1-15 (1987); Hillel *et al.*, *Anim. Genet.* 20:145-155 (1989); Hillel *et al.*, *Genet.* 124:783-789 (1990), all of which are herein incorporated by reference in their entirety).

The detection of polymorphic sites in a sample of DNA may be facilitated through the use of nucleic acid amplification methods. Such methods specifically increase the concentration of polynucleotides that span the polymorphic site, or include that site and sequences located either distal or proximal to it. Such amplified molecules can be readily detected by gel electrophoresis or other means.

The most preferred method of achieving such amplification employs the polymerase chain reaction ("PCR") (Mullis *et al.*, *Cold Spring Harbor Symp. Quant. Biol.* 51:263-273 (1986); Erlich *et al.*, European Patent Appln. 50,424; European Patent Appln. 84,796; European

Patent Application 258,017; European Patent Appln. 237,362; Mullis, European Patent Appln. 201,184; Mullis *et al.*, U.S. Patent No. 4,683,202; Erlich, U.S. Patent No. 4,582,788; and Saiki *et al.*, U.S. Patent No. 4,683,194), using primer pairs that are capable of hybridizing to the proximal sequences that define a polymorphism in its double-stranded form.

In lieu of PCR, alternative methods, such as the "Ligase Chain Reaction" ("LCR") may be used (Barany, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:189-193 (1991), the entirety of which is herein incorporated by reference). LCR uses two pairs of oligonucleotide probes to exponentially amplify a specific target. The sequences of each pair of oligonucleotides is selected to permit the pair to hybridize to abutting sequences of the same strand of the target. Such hybridization forms a substrate for a template-dependent ligase. As with PCR, the resulting products thus serve as a template in subsequent cycles and an exponential amplification of the desired sequence is obtained.

LCR can be performed with oligonucleotides having the proximal and distal sequences of the same strand of a polymorphic site. In one embodiment, either oligonucleotide will be designed to include the actual polymorphic site of the polymorphism. In such an embodiment, the reaction conditions are selected such that the oligonucleotides can be ligated together only if the target molecule either contains or lacks the specific nucleotide that is complementary to the polymorphic site present on the oligonucleotide. Alternatively, the oligonucleotides may be selected such that they do not include the polymorphic site (see, Segev, PCT Application WO 90/01069, the entirety of which is herein incorporated by reference).

The "Oligonucleotide Ligation Assay" ("OLA") may alternatively be employed (Landegren *et al.*, *Science* 241:1077-1080 (1988), the entirety of which is herein incorporated by reference). The OLA protocol uses two oligonucleotides which are designed to be capable of

hybridizing to abutting sequences of a single strand of a target. OLA, like LCR, is particularly suited for the detection of point mutations. Unlike LCR, however, OLA results in "linear" rather than exponential amplification of the target sequence.

Nickerson *et al.*, have described a nucleic acid detection assay that combines attributes of PCR and OLA (Nickerson *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:8923-8927 (1990), the entirety of which is herein incorporated by reference). In this method, PCR is used to achieve the exponential amplification of target DNA, which is then detected using OLA. In addition to requiring multiple and separate, processing steps, one problem associated with such combinations is that they inherit all of the problems associated with PCR and OLA.

Schemes based on ligation of two (or more) oligonucleotides in the presence of nucleic acid having the sequence of the resulting "di-oligonucleotide", thereby amplifying the di-oligonucleotide, are also known (Wu *et al.*, *Genomics* 4:560-569 (1989), the entirety of which is herein incorporated by reference) and may be readily adapted to the purposes of the present invention.

Other known nucleic acid amplification procedures, such as allele-specific oligomers, branched DNA technology, transcription-based amplification systems, or isothermal amplification methods may also be used to amplify and analyze such polymorphisms (Malek *et al.*, U.S. Patent 5,130,238; Davey *et al.*, European Patent Application 329,822; Schuster *et al.*, U.S. Patent 5,169,766; Miller *et al.*, PCT Patent Application WO 89/06700; Kwoh *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:1173-1177 (1989); Gingeras *et al.*, PCT Patent Application WO 88/10315; Walker *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 89:392-396 (1992), all of which are herein incorporated by reference in their entirety).

The identification of a polymorphism can be determined in a variety of ways. By correlating the presence or absence of it in a plant with the presence or absence of a phenotype, it is possible to predict the phenotype of that plant. If a polymorphism creates or destroys a restriction endonuclease cleavage site, or if it results in the loss or insertion of DNA (e.g., a VNTR polymorphism), it will alter the size or profile of the DNA fragments that are generated by digestion with that restriction endonuclease. As such, individuals that possess a variant sequence can be distinguished from those having the original sequence by restriction fragment analysis. Polymorphisms that can be identified in this manner are termed “restriction fragment length polymorphisms” (“RFLPs”). RFLPs have been widely used in human and plant genetic analyses (Glassberg, UK Patent Application 2135774; Skolnick *et al.*, *Cytogen. Cell Genet.* 32:58-67 (1982); Botstein *et al.*, *Ann. J. Hum. Genet.* 32:314-331 (1980); Fischer *et al.*, (PCT Application WO90/13668); Uhlen, PCT Application WO90/11369).

Polymorphisms can also be identified by Single Strand Conformation Polymorphism (SSCP) analysis. SSCP is a method capable of identifying most sequence variations in a single strand of DNA, typically between 150 and 250 nucleotides in length (Elles, *Methods in Molecular Medicine: Molecular Diagnosis of Genetic Diseases*, Humana Press (1996), the entirety of which is herein incorporated by reference); Orita *et al.*, *Genomics* 5:874-879 (1989), the entirety of which is herein incorporated by reference). Under denaturing conditions a single strand of DNA will adopt a conformation that is uniquely dependent on its sequence conformation. This conformation usually will be different, even if only a single base is changed. Most conformations have been reported to alter the physical configuration or size sufficiently to be detectable by electrophoresis. A number of protocols have been described for SSCP including, but not limited to, Lee *et al.*, *Anal. Biochem.* 205:289-293 (1992), the entirety of

which is herein incorporated by reference; Suzuki *et al.*, *Anal. Biochem.* 192:82-84 (1991), the entirety of which is herein incorporated by reference; Lo *et al.*, *Nucleic Acids Research* 20:1005-1009 (1992), the entirety of which is herein incorporated by reference; Sarkar *et al.*, *Genomics* 13:441-443 (1992), the entirety of which is herein incorporated by reference. It is understood that one or more of the nucleic acids of the present invention, may be utilized as markers or probes to detect polymorphisms by SSCP analysis.

Polymorphisms may also be found using a DNA fingerprinting technique called amplified fragment length polymorphism (AFLP), which is based on the selective PCR amplification of restriction fragments from a total digest of genomic DNA to profile that DNA (Vos *et al.*, *Nucleic Acids Res.* 23:4407-4414 (1995), the entirety of which is herein incorporated by reference). This method allows for the specific co-amplification of high numbers of restriction fragments, which can be visualized by PCR without knowledge of the nucleic acid sequence.

AFLP employs basically three steps. Initially, a sample of genomic DNA is cut with restriction enzymes and oligonucleotide adapters are ligated to the restriction fragments of the DNA. The restriction fragments are then amplified using PCR by using the adapter and restriction sequence as target sites for primer annealing. The selective amplification is achieved by the use of primers that extend into the restriction fragments, amplifying only those fragments in which the primer extensions match the nucleotide flanking the restriction sites. These amplified fragments are then visualized on a denaturing polyacrylamide gel.

AFLP analysis has been performed on *Salix* (Beismann *et al.*, *Mol. Ecol.* 6:989-993 (1997), the entirety of which is herein incorporated by reference), *Acinetobacter* (Janssen *et al.*, *Int. J. Syst. Bacteriol.* 47:1179-1187 (1997), the entirety of which is herein incorporated by

reference), *Aeromonas popoffi* (Huys *et al.*, *Int. J. Syst. Bacteriol.* 47:1165-1171 (1997), the entirety of which is herein incorporated by reference), rice (McCouch *et al.*, *Plant Mol. Biol.* 35:89-99 (1997), the entirety of which is herein incorporated by reference; Nandi *et al.*, *Mol. Gen. Genet.* 255:1-8 (1997), the entirety of which is herein incorporated by reference; Cho *et al.*, *Genome* 39:373-378 (1996), the entirety of which is herein incorporated by reference), barley (*Hordeum vulgare*)(Simons *et al.*, *Genomics* 44:61-70 (1997), the entirety of which is herein incorporated by reference; Waugh *et al.*, *Mol. Gen. Genet.* 255:311-321 (1997), the entirety of which is herein incorporated by reference; Qi *et al.*, *Mol. Gen. Genet.* 254:330-336 (1997), the entirety of which is herein incorporated by reference; Becker *et al.*, *Mol. Gen. Genet.* 249:65-73 (1995), the entirety of which is herein incorporated by reference), potato (Van der Voort *et al.*, *Mol. Gen. Genet.* 255:438-447 (1997), the entirety of which is herein incorporated by reference; Meksem *et al.*, *Mol. Gen. Genet.* 249:74-81 (1995), the entirety of which is herein incorporated by reference), *Phytophthora infestans* (Van der Lee *et al.*, *Fungal Genet. Biol.* 21:278-291 (1997), the entirety of which is herein incorporated by reference), *Bacillus anthracis* (Keim *et al.*, *J. Bacteriol.* 179:818-824 (1997), the entirety of which is herein incorporated by reference), *Astragalus cremnophylax* (Travis *et al.*, *Mol. Ecol.* 5:735-745 (1996), the entirety of which is herein incorporated by reference), *Arabidopsis* (Cnops *et al.*, *Mol. Gen. Genet.* 253:32-41 (1996), the entirety of which is herein incorporated by reference), *Escherichia coli* (Lin *et al.*, *Nucleic Acids Res.* 24:3649-3650 (1996), the entirety of which is herein incorporated by reference), *Aeromonas* (Huys *et al.*, *Int. J. Syst. Bacteriol.* 46:572-580 (1996), the entirety of which is herein incorporated by reference), nematode (Folkertsma *et al.*, *Mol. Plant Microbe Interact.* 9:47-54 (1996), the entirety of which is herein incorporated by reference), tomato (Thomas *et al.*, *Plant J.* 8:785-794 (1995), the entirety of which is herein incorporated by reference) and human (Latorra

*et al.*, *PCR Methods Appl.* 3:351-358 (1994), the entirety of which is herein incorporated by reference). AFLP analysis has also been used for fingerprinting mRNA (Money *et al.*, *Nucleic Acids Res.* 24:2616-2617 (1996), the entirety of which is herein incorporated by reference; Bachem *et al.*, *Plant J.* 9:745-753 (1996), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acids of the present invention, may be utilized as markers or probes to detect polymorphisms by AFLP analysis or for fingerprinting RNA.

Polymorphisms may also be found using random amplified polymorphic DNA (RAPD) (Williams *et al.*, *Nucl. Acids Res.* 18:6531-6535 (1990), the entirety of which is herein incorporated by reference) and cleaveable amplified polymorphic sequences (CAPS) (Lyamichev *et al.*, *Science* 260:778-783 (1993), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acid molecules of the present invention, may be utilized as markers or probes to detect polymorphisms by RAPD or CAPS analysis.

Through genetic mapping, a fine scale linkage map can be developed using DNA markers and, then, a genomic DNA library of large-sized fragments can be screened with molecular markers linked to the desired trait. Molecular markers are advantageous for agronomic traits that are otherwise difficult to tag, such as resistance to pathogens, insects and nematodes, tolerance to abiotic stress, quality parameters and quantitative traits such as high yield potential.

The essential requirements for marker-assisted selection in a plant breeding program are: (1) the marker(s) should co-segregate or be closely linked with the desired trait; (2) an efficient means of screening large populations for the molecular marker(s) should be available; and (3) the screening technique should have high reproducibility across laboratories and preferably be economical to use and be user-friendly.

The genetic linkage of marker molecules can be established by a gene mapping model such as, without limitation, the flanking marker model reported by Lander and Botstein, *Genetics* 121:185-199 (1989) and the interval mapping, based on maximum likelihood methods described by Lander and Botstein, *Genetics* 121:185-199 (1989) and implemented in the software package MAPMAKER/QTL (Lincoln and Lander, *Mapping Genes Controlling Quantitative Traits Using MAPMAKER/QTL*, Whitehead Institute for Biomedical Research, Massachusetts, (1990). Additional software includes Qgene, Version 2.23 (1996), Department of Plant Breeding and Biometry, 266 Emerson Hall, Cornell University, Ithaca, NY, the manual of which is herein incorporated by reference in its entirety). Use of Qgene software is a particularly preferred approach.

A maximum likelihood estimate (MLE) for the presence of a marker is calculated, together with an MLE assuming no QTL effect, to avoid false positives. A  $\log_{10}$  of an odds ratio (LOD) is then calculated as:  $LOD = \log_{10} (MLE \text{ for the presence of a QTL} / MLE \text{ given no linked QTL})$ .

The LOD score essentially indicates how much more likely the data are to have arisen assuming the presence of a QTL than in its absence. The LOD threshold value for avoiding a false positive with a given confidence, say 95%, depends on the number of markers and the length of the genome. Graphs indicating LOD thresholds are set forth in Lander and Botstein, *Genetics* 121:185-199 (1989) the entirety of which is herein incorporated by reference and further described by Arús and Moreno-González, *Plant Breeding*, Hayward *et al.*, (eds.) Chapman & Hall, London, pp. 314-331 (1993), the entirety of which is herein incorporated by reference.



Additional models can be used. Many modifications and alternative approaches to interval mapping have been reported, including the use non-parametric methods (Kruglyak and Lander, *Genetics* 139:1421-1428 (1995), the entirety of which is herein incorporated by reference). Multiple regression methods or models can be also be used, in which the trait is regressed on a large number of markers (Jansen, *Biometrics in Plant Breeding*, van Oijen and Jansen (eds.), Proceedings of the Ninth Meeting of the Eucarpia Section Biometrics in Plant Breeding, The Netherlands, pp. 116-124 (1994); Weber and Wricke, *Advances in Plant Breeding*, Blackwell, Berlin, 16 (1994), both of which is herein incorporated by reference in their entirety).

Procedures combining interval mapping with regression analysis, whereby the phenotype is regressed onto a single putative QTL at a given marker interval and at the same time onto a number of markers that serve as 'cofactors,' have been reported by Jansen and Stam, *Genetics* 136:1447-1455 (1994), the entirety of which is herein incorporated by reference and Zeng, *Genetics* 136:1457-1468 (1994) the entirety of which is herein incorporated by reference. Generally, the use of cofactors reduces the bias and sampling error of the estimated QTL positions (Utz and Melchinger, *Biometrics in Plant Breeding*, van Oijen and Jansen (eds.) Proceedings of the Ninth Meeting of the Eucarpia Section Biometrics in Plant Breeding, The Netherlands, pp.195-204 (1994), the entirety of which is herein incorporated by reference, thereby improving the precision and efficiency of QTL mapping (Zeng, *Genetics* 136:1457-1468 (1994)). These models can be extended to multi-environment experiments to analyze genotype-environment interactions (Jansen *et al.*, *Theo. Appl. Genet.* 91:33-37 (1995), the entirety of which is herein incorporated by reference).

Selection of an appropriate mapping populations is important to map construction. The choice of appropriate mapping population depends on the type of marker systems employed

(Tanksley *et al.*, *Molecular mapping plant chromosomes. Chromosome structure and function: Impact of new concepts*, Gustafson and Appels (eds.), Plenum Press, New York, pp 157-173 (1988), the entirety of which is herein incorporated by reference). Consideration must be given to the source of parents (adapted vs. exotic) used in the mapping population. Chromosome pairing and recombination rates can be severely disturbed (suppressed) in wide crosses (adapted x exotic) and generally yield greatly reduced linkage distances. Wide crosses will usually provide segregating populations with a relatively large array of polymorphisms when compared to progeny in a narrow cross (adapted x adapted).

An  $F_2$  population is the first generation of selfing after the hybrid seed is produced. Usually a single  $F_1$  plant is selfed to generate a population segregating for all the genes in Mendelian (1:2:1) fashion. Maximum genetic information is obtained from a completely classified  $F_2$  population using a codominant marker system (Mather, *Measurement of Linkage in Heredity*, Methuen and Co., (1938), the entirety of which is herein incorporated by reference). In the case of dominant markers, progeny tests (e.g.  $F_3$ ,  $BCF_2$ ) are required to identify the heterozygotes, thus making it equivalent to a completely classified  $F_2$  population. However, this procedure is often prohibitive because of the cost and time involved in progeny testing. Progeny testing of  $F_2$  individuals is often used in map construction where phenotypes do not consistently reflect genotype (e.g. disease resistance) or where trait expression is controlled by a QTL. Segregation data from progeny test populations (e.g.  $F_3$  or  $BCF_2$ ) can be used in map construction. Marker-assisted selection can then be applied to cross progeny based on marker-trait map associations ( $F_2$ ,  $F_3$ ), where linkage groups have not been completely disassociated by recombination events (i.e., maximum disequilibrium).

Recombinant inbred lines (RIL) (genetically related lines; usually  $>F_5$ , developed from continuously selfing  $F_2$  lines towards homozygosity) can be used as a mapping population. Information obtained from dominant markers can be maximized by using RIL because all loci are homozygous or nearly so. Under conditions of tight linkage (i.e., about  $<10\%$  recombination), dominant and co-dominant markers evaluated in RIL populations provide more information per individual than either marker type in backcross populations (Reiter *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 89:1477-1481 (1992), the entirety of which is herein incorporated by reference). However, as the distance between markers becomes larger (i.e., loci become more independent), the information in RIL populations decreases dramatically when compared to codominant markers.

Backcross populations (e.g., generated from a cross between a successful variety (recurrent parent) and another variety (donor parent) carrying a trait not present in the former) can be utilized as a mapping population. A series of backcrosses to the recurrent parent can be made to recover most of its desirable traits. Thus a population is created consisting of individuals nearly like the recurrent parent but each individual carries varying amounts or mosaic of genomic regions from the donor parent. Backcross populations can be useful for mapping dominant markers if all loci in the recurrent parent are homozygous and the donor and recurrent parent have contrasting polymorphic marker alleles (Reiter *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 89:1477-1481 (1992)). Information obtained from backcross populations using either codominant or dominant markers is less than that obtained from  $F_2$  populations because one, rather than two, recombinant gametes are sampled per plant. Backcross populations, however, are more informative (at low marker saturation) when compared to RILs as the distance between linked loci increases in RIL populations (i.e. about  $15\%$  recombination). Increased

recombination can be beneficial for resolution of tight linkages, but may be undesirable in the construction of maps with low marker saturation.

Near-isogenic lines (NIL) created by many backcrosses to produce an array of individuals that are nearly identical in genetic composition except for the trait or genomic region under interrogation can be used as a mapping population. In mapping with NILs, only a portion of the polymorphic loci are expected to map to a selected region.

Bulk segregant analysis (BSA) is a method developed for the rapid identification of linkage between markers and traits of interest (Michelmore *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:9828-9832 (1991), the entirety of which is herein incorporated by reference). In BSA, two bulked DNA samples are drawn from a segregating population originating from a single cross. These bulks contain individuals that are identical for a particular trait (resistant or susceptible to particular disease) or genomic region but arbitrary at unlinked regions (i.e. heterozygous). Regions unlinked to the target region will not differ between the bulked samples of many individuals in BSA.

It is understood that one or more of the nucleic acid molecules of the present invention may be used as molecular markers. It is also understood that one or more of the protein molecules of the present invention may be used as molecular markers.

In accordance with this aspect of the present invention, a sample nucleic acid is obtained from plants cells or tissues. Any source of nucleic acid may be used. Preferably, the nucleic acid is genomic DNA. The nucleic acid is subjected to restriction endonuclease digestion. For example, one or more nucleic acid molecule or fragment thereof of the present invention can be used as a probe in accordance with the above-described polymorphic methods. The polymorphism obtained in this approach can then be cloned to identify the mutation at the coding

region which alters the protein's structure or regulatory region of the gene which affects its expression level.

In an aspect of the present invention, one or more of the nucleic molecules of the present invention are used to determine the level (i.e., the concentration of mRNA in a sample, etc.) in a plant (preferably maize or soybean) or pattern (i.e., the kinetics of expression, rate of decomposition, stability profile, etc.) of the expression of a protein encoded in part or whole by one or more of the nucleic acid molecule of the present invention (collectively, the "Expression Response" of a cell or tissue). As used herein, the Expression Response manifested by a cell or tissue is said to be "altered" if it differs from the Expression Response of cells or tissues of plants not exhibiting the phenotype. To determine whether a Expression Response is altered, the Expression Response manifested by the cell or tissue of the plant exhibiting the phenotype is compared with that of a similar cell or tissue sample of a plant not exhibiting the phenotype. As will be appreciated, it is not necessary to re-determine the Expression Response of the cell or tissue sample of plants not exhibiting the phenotype each time such a comparison is made; rather, the Expression Response of a particular plant may be compared with previously obtained values of normal plants. As used herein, the phenotype of the organism is any of one or more characteristics of an organism (e.g. disease resistance, pest tolerance, environmental tolerance such as tolerance to abiotic stress, male sterility, quality improvement or yield etc.). A change in genotype or phenotype may be transient or permanent. Also as used herein, a tissue sample is any sample that comprises more than one cell. In a preferred aspect, a tissue sample comprises cells that share a common characteristic (e.g. derived from root, seed, flower, leaf, stem or pollen etc.).

In one aspect of the present invention, an evaluation can be conducted to determine whether a particular mRNA molecule is present. One or more of the nucleic acid molecules of the present invention, preferably one or more of the EST nucleic acid molecules or fragments thereof of the present invention are utilized to detect the presence or quantity of the mRNA species. Such molecules are then incubated with cell or tissue extracts of a plant under conditions sufficient to permit nucleic acid hybridization. The detection of double-stranded probe-mRNA hybrid molecules is indicative of the presence of the mRNA; the amount of such hybrid formed is proportional to the amount of mRNA. Thus, such probes may be used to ascertain the level and extent of the mRNA production in a plant's cells or tissues. Such nucleic acid hybridization may be conducted under quantitative conditions (thereby providing a numerical value of the amount of the mRNA present). Alternatively, the assay may be conducted as a qualitative assay that indicates either that the mRNA is present, or that its level exceeds a user set, predefined value.

A principle of *in situ* hybridization is that a labeled, single-stranded nucleic acid probe will hybridize to a complementary strand of cellular DNA or RNA and, under the appropriate conditions, these molecules will form a stable hybrid. When nucleic acid hybridization is combined with histological techniques, specific DNA or RNA sequences can be identified within a single cell. An advantage of *in situ* hybridization over more conventional techniques for the detection of nucleic acids is that it allows an investigator to determine the precise spatial population (Angerer *et al.*, *Dev. Biol.* 101:477-484 (1984), the entirety of which is herein incorporated by reference; Angerer *et al.*, *Dev. Biol.* 112:157-166 (1985), the entirety of which is herein incorporated by reference; Dixon *et al.*, *EMBO J.* 10:1317-1324 (1991), the entirety of which is herein incorporated by reference). *In situ* hybridization may be used to measure the

steady-state level of RNA accumulation. It is a sensitive technique and RNA sequences present in as few as 5-10 copies per cell can be detected (Hardin *et al.*, *J. Mol. Biol.* 202:417-431 (1989), the entirety of which is herein incorporated by reference). A number of protocols have been devised for *in situ* hybridization, each with tissue preparation, hybridization and washing conditions (Meyerowitz, *Plant Mol. Biol. Rep.* 5:242-250 (1987), the entirety of which is herein incorporated by reference; Cox and Goldberg, In: *Plant Molecular Biology: A Practical Approach*, Shaw (ed.), pp 1-35, IRL Press, Oxford (1988), the entirety of which is herein incorporated by reference; Raikhel *et al.*, *In situ RNA hybridization in plant tissues*, In: *Plant Molecular Biology Manual*, vol. B9:1-32, Kluwer Academic Publisher, Dordrecht, Belgium (1989), the entirety of which is herein incorporated by reference).

*In situ* hybridization also allows for the localization of proteins within a tissue or cell (Wilkinson, *In Situ Hybridization*, Oxford University Press, Oxford (1992), the entirety of which is herein incorporated by reference; Langdale, *In Situ Hybridization* In: *The Maize Handbook*, Freeling and Walbot (eds.), pp 165-179, Springer-Verlag, New York (1994), the entirety of which is herein incorporated by reference). It is understood that one or more of the molecules of the present invention, preferably one or more of the EST nucleic acid molecules or fragments thereof of the present invention or one or more of the antibodies of the present invention may be utilized to detect the level or pattern of a methionine pathway protein or mRNA thereof by *in situ* hybridization.

Fluorescent *in situ* hybridization allows the localization of a particular DNA sequence along a chromosome which is useful, among other uses, for gene mapping, following chromosomes in hybrid lines or detecting chromosomes with translocations, transversions or deletions. *In situ* hybridization has been used to identify chromosomes in several plant species

(Griffor *et al.*, *Plant Mol. Biol.* 17:101-109 (1991), the entirety of which is herein incorporated by reference; Gustafson *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:1899-1902 (1990), herein incorporated by reference; Mukai and Gill, *Genome* 34:448-452 (1991), the entirety of which is herein incorporated by reference; Schwarzach and Heslop-Harrison, *Genome* 34:317-323 (1991); Wang *et al.*, *Jpn. J. Genet.* 66:313-316 (1991), the entirety of which is herein incorporated by reference; Parra and Windle, *Nature Genetics* 5:17-21 (1993), the entirety of which is herein incorporated by reference). It is understood that the nucleic acid molecules of the present invention may be used as probes or markers to localize sequences along a chromosome.

Another method to localize the expression of a molecule is tissue printing. Tissue printing provides a way to screen, at the same time on the same membrane many tissue sections from different plants or different developmental stages. Tissue-printing procedures utilize films designed to immobilize proteins and nucleic acids. In essence, a freshly cut section of a tissue is pressed gently onto nitrocellulose paper, nylon membrane or polyvinylidene difluoride membrane. Such membranes are commercially available (*e.g.* Millipore, Bedford, Massachusetts U.S.A.). The contents of the cut cell transfer onto the membrane and the contents and are immobilized to the membrane. The immobilized contents form a latent print that can be visualized with appropriate probes. When a plant tissue print is made on nitrocellulose paper, the cell walls leave a physical print that makes the anatomy visible without further treatment (Varner and Taylor, *Plant Physiol.* 91:31-33 (1989), the entirety of which is herein incorporated by reference).

Tissue printing on substrate films is described by Daoust, *Exp. Cell Res.* 12:203-211 (1957), the entirety of which is herein incorporated by reference, who detected amylase, protease, ribonuclease and deoxyribonuclease in animal tissues using starch, gelatin and agar films. These



techniques can be applied to plant tissues (Yomo and Taylor, *Planta* 112:35-43 (1973); the entirety of which is herein incorporated by reference; Harris and Chrispeels, *Plant Physiol.* 56:292-299 (1975), the entirety of which is herein incorporated by reference). Advances in membrane technology have increased the range of applications of Daoust's tissue-printing techniques allowing (Cassab and Varner, *J. Cell. Biol.* 105:2581-2588 (1987), the entirety of which is herein incorporated by reference) the histochemical localization of various plant enzymes and deoxyribonuclease on nitrocellulose paper and nylon (Spruce *et al.*, *Phytochemistry* 26:2901-2903 (1987), the entirety of which is herein incorporated by reference; Barres *et al.*, *Neuron* 5:527-544 (1990), the entirety of which is herein incorporated by reference; Reid and Pont-Lezica, *Tissue Printing: Tools for the Study of Anatomy, Histochemistry and Gene Expression*, Academic Press, New York, New York (1992), the entirety of which is herein incorporated by reference; Reid *et al.*, *Plant Physiol.* 93:160-165 (1990), the entirety of which is herein incorporated by reference; Ye *et al.*, *Plant J.* 1:175-183 (1991), the entirety of which is herein incorporated by reference).

It is understood that one or more of the molecules of the present invention, preferably one or more of the EST nucleic acid molecules or fragments thereof of the present invention or one or more of the antibodies of the present invention may be utilized to detect the presence or quantity of a methionine pathway protein by tissue printing.

Further it is also understood that any of the nucleic acid molecules of the present invention may be used as marker nucleic acids and or probes in connection with methods that require probes or marker nucleic acids. As used herein, a probe is an agent that is utilized to determine an attribute or feature (e.g. presence or absence, location, correlation, etc.) of a molecule, cell, tissue or plant. As used herein, a marker nucleic acid is a nucleic acid molecule

that is utilized to determine an attribute or feature (*e.g.*, presence or absence, location, correlation, etc.) or a molecule, cell, tissue or plant.

A microarray-based method for high-throughput monitoring of plant gene expression may be utilized to measure gene-specific hybridization targets. This 'chip'-based approach involves using microarrays of nucleic acid molecules as gene-specific hybridization targets to quantitatively measure expression of the corresponding plant genes (Schena *et al.*, *Science* 270:467-470 (1995), the entirety of which is herein incorporated by reference; Shalon, Ph.D. Thesis, Stanford University (1996), the entirety of which is herein incorporated by reference). Every nucleotide in a large sequence can be queried at the same time. Hybridization can be used to efficiently analyze nucleotide sequences.

Several microarray methods have been described. One method compares the sequences to be analyzed by hybridization to a set of oligonucleotides representing all possible subsequences (Bains and Smith, *J. Theor. Biol.* 135:303-307 (1989), the entirety of which is herein incorporated by reference). A second method hybridizes the sample to an array of oligonucleotide or cDNA molecules. An array consisting of oligonucleotides complementary to subsequences of a target sequence can be used to determine the identity of a target sequence, measure its amount and detect differences between the target and a reference sequence. Nucleic acid molecules microarrays may also be screened with protein molecules or fragments thereof to determine nucleic acid molecules that specifically bind protein molecules or fragments thereof.

The microarray approach may be used with polypeptide targets (U.S. Patent No. 5,445,934; U.S. Patent No. 5,143,854; U.S. Patent No. 5,079,600; U.S. Patent No. 4,923,901, all of which are herein incorporated by reference in their entirety). Essentially, polypeptides are synthesized on a substrate (microarray) and these polypeptides can be screened with either

protein molecules or fragments thereof or nucleic acid molecules in order to screen for either protein molecules or fragments thereof or nucleic acid molecules that specifically bind the target polypeptides. (Fodor *et al.*, *Science* 251:767-773 (1991), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acid molecules or protein or fragments thereof of the present invention may be utilized in a microarray based method.

In a preferred embodiment of the present invention microarrays may be prepared that comprise nucleic acid molecules where such nucleic acid molecules encode at least one, preferably at least two, more preferably at least three, even more preferably at least four, five six or seven methionine pathway enzymes. In a preferred embodiment the nucleic acid molecules are selected from the group consisting of a nucleic acid molecule that encodes a maize or a soybean methionine adenosyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean S-adenosylmethionine decarboxylase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate kinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean aspartate-semialdehyde dehydrogenase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean *O*-succinylhomoserine (thiol)-lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -lyase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or fragment thereof, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or fragment thereof, a nucleic acid molecule that encodes a

maize or a soybean cystathionine  $\gamma$ -lyase enzyme or fragment thereof and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or fragment thereof.

Site directed mutagenesis may be utilized to modify nucleic acid sequences, particularly as it is a technique that allows one or more of the amino acids encoded by a nucleic acid molecule to be altered (e.g. a threonine to be replaced by a methionine). Three basic methods for site directed mutagenesis are often employed. These are cassette mutagenesis (Wells *et al.*, *Gene* 34:315-323 (1985), the entirety of which is herein incorporated by reference), primer extension (Gilliam *et al.*, *Gene* 12:129-137 (1980), the entirety of which is herein incorporated by reference; Zoller and Smith, *Methods Enzymol.* 100:468-500 (1983), the entirety of which is herein incorporated by reference; Dalbadie-McFarland *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 79:6409-6413 (1982), the entirety of which is herein incorporated by reference) and methods based upon PCR (Scharf *et al.*, *Science* 233:1076-1078 (1986), the entirety of which is herein incorporated by reference; Higuchi *et al.*, *Nucleic Acids Res.* 16:7351-7367 (1988), the entirety of which is herein incorporated by reference). Site directed mutagenesis approaches are also described in European Patent 0 385 962, the entirety of which is herein incorporated by reference; European Patent 0 359 472, the entirety of which is herein incorporated by reference; and PCT Patent Application WO 93/07278, the entirety of which is herein incorporated by reference.

Site directed mutagenesis strategies have been applied to plants for both *in vitro* as well as *in vivo* site directed mutagenesis (Lanz *et al.*, *J. Biol. Chem.* 266:9971-9976 (1991), the entirety of which is herein incorporated by reference; Kovgan and Zhdanov, *Biotekhnologiya* 5:148-154, No. 207160n, Chemical Abstracts 110:225 (1989), the entirety of which is herein incorporated by reference; Ge *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:4037-4041 (1989), the

entirety of which is herein incorporated by reference; Zhu *et al.*, *J. Biol. Chem.* 271:18494-18498 (1996), the entirety of which is herein incorporated by reference; Chu *et al.*, *Biochemistry* 33:6150-6157 (1994), the entirety of which is herein incorporated by reference; Small *et al.*, *EMBO J.* 11:1291-1296 (1992), the entirety of which is herein incorporated by reference; Cho *et al.*, *Mol. Biotechnol.* 8:13-16 (1997), the entirety of which is herein incorporated by reference; Kita *et al.*, *J. Biol. Chem.* 271:26529-26535 (1996), the entirety of which is herein incorporated by reference; Jin *et al.*, *Mol. Microbiol.* 7:555-562 (1993), the entirety of which is herein incorporated by reference; Hatfield and Vierstra, *J. Biol. Chem.* 267:14799-14803 (1992), the entirety of which is herein incorporated by reference; Zhao *et al.*, *Biochemistry* 31:5093-5099 (1992), the entirety of which is herein incorporated by reference).

Any of the nucleic acid molecules of the present invention may either be modified by site directed mutagenesis or used as, for example, nucleic acid molecules that are used to target other nucleic acid molecules for modification. It is understood that mutants with more than one altered nucleotide can be constructed using techniques that practitioners are familiar with such as isolating restriction fragments and ligating such fragments into an expression vector (*see, for example, Sambrook et al., Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Press (1989)).

Sequence-specific DNA-binding proteins play a role in the regulation of transcription. The isolation of recombinant cDNAs encoding these proteins facilitates the biochemical analysis of their structural and functional properties. Genes encoding such DNA-binding proteins have been isolated using classical genetics (Vollbrecht *et al.*, *Nature* 350: 241-243 (1991), the entirety of which is herein incorporated by reference) and molecular biochemical approaches, including the screening of recombinant cDNA libraries with antibodies (Landschulz *et al.*, *Genes Dev.*

2:786-800 (1988), the entirety of which is herein incorporated by reference) or DNA probes (Bodner *et al.*, *Cell* 55:505-518 (1988), the entirety of which is herein incorporated by reference). In addition, an *in situ* screening procedure has been used and has facilitated the isolation of sequence-specific DNA-binding proteins from various plant species (Gilmartin *et al.*, *Plant Cell* 4:839-849 (1992), the entirety of which is herein incorporated by reference; Schindler *et al.*, *EMBO J.* 11:1261-1273 (1992), the entirety of which is herein incorporated by reference). An *in situ* screening protocol does not require the purification of the protein of interest (Vinson *et al.*, *Genes Dev.* 2:801-806 (1988), the entirety of which is herein incorporated by reference; Singh *et al.*, *Cell* 52:415-423 (1988), the entirety of which is herein incorporated by reference).

Two steps may be employed to characterize DNA-protein interactions. The first is to identify promoter fragments that interact with DNA-binding proteins, to titrate binding activity, to determine the specificity of binding and to determine whether a given DNA-binding activity can interact with related DNA sequences (Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, 2<sup>nd</sup> edition, Cold Spring Harbor Laboratory Press, Cold Spring Harbor, New York (1989)). Electrophoretic mobility-shift assay is a widely used assay. The assay provides a rapid and sensitive method for detecting DNA-binding proteins based on the observation that the mobility of a DNA fragment through a nondenaturing, low-ionic strength polyacrylamide gel is retarded upon association with a DNA-binding protein (Fried and Crother, *Nucleic Acids Res.* 9:6505-6525 (1981), the entirety of which is herein incorporated by reference). When one or more specific binding activities have been identified, the exact sequence of the DNA bound by the protein may be determined. Several procedures for characterizing protein/DNA-binding sites are used, including methylation and ethylation interference assays (Maxam and Gilbert, *Methods*

*Enzymol.* 65:499-560 (1980), the entirety of which is herein incorporated by reference; Wissman and Hillen, *Methods Enzymol.* 208:365-379 (1991), the entirety of which is herein incorporated by reference), footprinting techniques employing DNase I (Galas and Schmitz, *Nucleic Acids Res.* 5:3157-3170 (1978), the entirety of which is herein incorporated by reference), 1,10-phenanthroline-copper ion methods (Sigman *et al.*, *Methods Enzymol.* 208:414-433 (1991), the entirety of which is herein incorporated by reference) and hydroxyl radicals methods (Dixon *et al.*, *Methods Enzymol.* 208:414-433 (1991), the entirety of which is herein incorporated by reference). It is understood that one or more of the nucleic acid molecules of the present invention may be utilized to identify a protein or fragment thereof that specifically binds to a nucleic acid molecule of the present invention. It is also understood that one or more of the protein molecules or fragments thereof of the present invention may be utilized to identify a nucleic acid molecule that specifically binds to it.

A two-hybrid system is based on the fact that many cellular functions are carried out by proteins, such as transcription factors, that interact (physically) with one another. Two-hybrid systems have been used to probe the function of new proteins (Chien *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 88:9578-9582 (1991) the entirety of which is herein incorporated by reference; Durfee *et al.*, *Genes Dev.* 7:555-569 (1993) the entirety of which is herein incorporated by reference; Choi *et al.*, *Cell* 78:499-512 (1994), the entirety of which is herein incorporated by reference; Kranz *et al.*, *Genes Dev.* 8:313-327 (1994), the entirety of which is herein incorporated by reference).

Interaction mating techniques have facilitated a number of two-hybrid studies of protein-protein interaction. Interaction mating has been used to examine interactions between small sets of tens of proteins (Finley and Brent, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:12098-12984 (1994), the entirety of which is herein incorporated by reference), larger sets of hundreds of proteins

(Bendixen *et al.*, *Nucl. Acids Res.* 22:1778-1779 (1994), the entirety of which is herein incorporated by reference) and to comprehensively map proteins encoded by a small genome (Bartel *et al.*, *Nature Genetics* 12:72-77 (1996), the entirety of which is herein incorporated by reference). This technique utilizes proteins fused to the DNA-binding domain and proteins fused to the activation domain. They are expressed in two different haploid yeast strains of opposite mating type and the strains are mated to determine if the two proteins interact. Mating occurs when haploid yeast strains come into contact and result in the fusion of the two haploids into a diploid yeast strain. An interaction can be determined by the activation of a two-hybrid reporter gene in the diploid strain. An advantage of this technique is that it reduces the number of yeast transformations needed to test individual interactions. It is understood that the protein-protein interactions of protein or fragments thereof of the present invention may be investigated using the two-hybrid system and that any of the nucleic acid molecules of the present invention that encode such proteins or fragments thereof may be used to transform yeast in the two-hybrid system.

#### **(a) Plant Constructs and Plant Transformants**

One or more of the nucleic acid molecules of the present invention may be used in plant transformation or transfection. Exogenous genetic material may be transferred into a plant cell and the plant cell regenerated into a whole, fertile or sterile plant. Exogenous genetic material is any genetic material, whether naturally occurring or otherwise, from any source that is capable of being inserted into any organism. Such genetic material may be transferred into either monocotyledons and dicotyledons including, but not limited to maize (pp 63-69), soybean (pp



50-60), *Arabidopsis* (p 45), phaseolus (pp 47-49), peanut (pp 49-50), alfalfa (p 60), wheat (pp 69-71), rice (pp 72-79), oat (pp 80-81), sorghum (p 83), rye (p 84), tritordeum (p 84), millet (p85), fescue (p 85), perennial ryegrass (p 86), sugarcane (p87), cranberry (p101), papaya (pp 101-102), banana (p 103), banana (p 103), muskmelon (p 104), apple (p 104), cucumber (p 105), dendrobium (p 109), gladiolus (p 110), chrysanthemum (p 110), liliacea (p 111), cotton (pp113-114), eucalyptus (p 115), sunflower (p 118), canola (p 118), turfgrass (p121), sugarbeet (p 122), coffee (p 122) and dioscorea (p 122), (Christou, In: *Particle Bombardment for Genetic Engineering of Plants*, Biotechnology Intelligence Unit. Academic Press, San Diego, California (1996), the entirety of which is herein incorporated by reference).

Transfer of a nucleic acid that encodes for a protein can result in overexpression of that protein in a transformed cell or transgenic plant. One or more of the proteins or fragments thereof encoded by nucleic acid molecules of the present invention may be overexpressed in a transformed cell or transformed plant. Particularly, any of the methionine pathway proteins or fragments thereof may be overexpressed in a transformed cell or transgenic plant. Such overexpression may be the result of transient or stable transfer of the exogenous genetic material.

Exogenous genetic material may be transferred into a plant cell and the plant cell by the use of a DNA vector or construct designed for such a purpose. Design of such a vector is generally within the skill of the art (See, *Plant Molecular Biology: A Laboratory Manual*, Clark (ed.), Springer, New York (1997), the entirety of which is herein incorporated by reference).

A construct or vector may include a plant promoter to express the protein or protein fragment of choice. A number of promoters which are active in plant cells have been described in the literature. These include the nopaline synthase (NOS) promoter (Ebert *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 84:5745-5749 (1987), the entirety of which is herein incorporated by

reference), the octopine synthase (OCS) promoter (which are carried on tumor-inducing plasmids of *Agrobacterium tumefaciens*), the caulimovirus promoters such as the cauliflower mosaic virus (CaMV) 19S promoter (Lawton *et al.*, *Plant Mol. Biol.* 9:315-324 (1987), the entirety of which is herein incorporated by reference) and the CAMV 35S promoter (Odell *et al.*, *Nature* 313:810-812 (1985), the entirety of which is herein incorporated by reference), the figwort mosaic virus 35S-promoter, the light-inducible promoter from the small subunit of ribulose-1,5-bis-phosphate carboxylase (ssRUBISCO), the Adh promoter (Walker *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 84:6624-6628 (1987), the entirety of which is herein incorporated by reference), the sucrose synthase promoter (Yang *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:4144-4148 (1990), the entirety of which is herein incorporated by reference), the R gene complex promoter (Chandler *et al.*, *The Plant Cell* 1:1175-1183 (1989), the entirety of which is herein incorporated by reference) and the chlorophyll a/b binding protein gene promoter, etc. These promoters have been used to create DNA constructs which have been expressed in plants; *see, e.g.*, PCT publication WO 84/02913, herein incorporated by reference in its entirety.

Promoters which are known or are found to cause transcription of DNA in plant cells can be used in the present invention. Such promoters may be obtained from a variety of sources such as plants and plant viruses. It is preferred that the particular promoter selected should be capable of causing sufficient expression to result in the production of an effective amount of the methionine pathway protein to cause the desired phenotype. In addition to promoters that are known to cause transcription of DNA in plant cells, other promoters may be identified for use in the current invention by screening a plant cDNA library for genes which are selectively or preferably expressed in the target tissues or cells.

For the purpose of expression in source tissues of the plant, such as the leaf, seed, root or stem, it is preferred that the promoters utilized in the present invention have relatively high expression in these specific tissues. For this purpose, one may choose from a number of promoters for genes with tissue- or cell-specific or -enhanced expression. Examples of such promoters reported in the literature include the chloroplast glutamine synthetase GS2 promoter from pea (Edwards *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:3459-3463 (1990), herein incorporated by reference in its entirety), the chloroplast fructose-1,6-biphosphatase (FBPase) promoter from wheat (Lloyd *et al.*, *Mol. Gen. Genet.* 225:209-216 (1991), herein incorporated by reference in its entirety), the nuclear photosynthetic ST-LS1 promoter from potato (Stockhaus *et al.*, *EMBO J.* 8:2445-2451 (1989), herein incorporated by reference in its entirety), the serine/threonine kinase (PAL) promoter and the glucoamylase (CHS) promoter from *Arabidopsis thaliana*. Also reported to be active in photosynthetically active tissues are the ribulose-1,5-bisphosphate carboxylase (RbcS) promoter from eastern larch (*Larix laricina*), the promoter for the *cab* gene, *cab6*, from pine (Yamamoto *et al.*, *Plant Cell Physiol.* 35:773-778 (1994), herein incorporated by reference in its entirety), the promoter for the *Cab-1* gene from wheat (Fejes *et al.*, *Plant Mol. Biol.* 15:921-932 (1990), herein incorporated by reference in its entirety), the promoter for the *CAB-1* gene from spinach (Lubberstedt *et al.*, *Plant Physiol.* 104:997-1006 (1994), herein incorporated by reference in its entirety), the promoter for the *cab1R* gene from rice (Luan *et al.*, *Plant Cell.* 4:971-981 (1992), the entirety of which is herein incorporated by reference), the pyruvate, orthophosphate dikinase (PPDK) promoter from maize (Matsuoka *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 90: 9586-9590 (1993), herein incorporated by reference in its entirety), the promoter for the tobacco *Lhcb1\*2* gene (Cerdan *et al.*, *Plant Mol. Biol.* 33:245-255 (1997), herein incorporated by reference in its entirety), the *Arabidopsis thaliana* SUC2 sucrose-

H<sup>+</sup> symporter promoter (Truernit *et al.*, *Planta*. 196:564-570 (1995), herein incorporated by reference in its entirety) and the promoter for the thylakoid membrane proteins from spinach (psaD, psaF, psaE, PC, FNR, atpC, atpD, cab, rbcS). Other promoters for the chlorophyll a/b-binding proteins may also be utilized in the present invention, such as the promoters for LhcB gene and PsbP gene from white mustard (*Sinapis alba*; Kretsch *et al.*, *Plant Mol. Biol.* 28:219-229 (1995), the entirety of which is herein incorporated by reference).

For the purpose of expression in sink tissues of the plant, such as the tuber of the potato plant, the fruit of tomato, or the seed of maize, wheat, rice and barley, it is preferred that the promoters utilized in the present invention have relatively high expression in these specific tissues. A number of promoters for genes with tuber-specific or -enhanced expression are known, including the class I patatin promoter (Bevan *et al.*, *EMBO J.* 8:1899-1906 (1986); Jefferson *et al.*, *Plant Mol. Biol.* 14:995-1006 (1990), both of which are herein incorporated by reference in its entirety), the promoter for the potato tuber ADPGPP genes, both the large and small subunits, the sucrose synthase promoter (Salanoubat and Belliard, *Gene*. 60:47-56 (1987), Salanoubat and Belliard, *Gene*. 84:181-185 (1989), both of which are incorporated by reference in their entirety), the promoter for the major tuber proteins including the 22 kd protein complexes and proteinase inhibitors (Hannapel, *Plant Physiol.* 101:703-704 (1993), herein incorporated by reference in its entirety), the promoter for the granule bound starch synthase gene (GBSS) (Visser *et al.*, *Plant Mol. Biol.* 17:691-699 (1991), herein incorporated by reference in its entirety) and other class I and II patatins promoters (Koster-Topfer *et al.*, *Mol Gen Genet.* 219:390-396 (1989); Mignery *et al.*, *Gene*. 62:27-44 (1988), both of which are herein incorporated by reference in their entirety).

Other promoters can also be used to express a methionine pathway protein or fragment thereof in specific tissues, such as seeds or fruits. The promoter for  $\beta$ -conglycinin (Chen *et al.*, *Dev. Genet.* 10: 112-122 (1989), herein incorporated by reference in its entirety) or other seed-specific promoters such as the napin and phaseolin promoters, can be used. The zeins are a group of storage proteins found in maize endosperm. Genomic clones for zein genes have been isolated (Pedersen *et al.*, *Cell* 29:1015-1026 (1982), herein incorporated by reference in its entirety) and the promoters from these clones, including the 15 kD, 16 kD, 19 kD, 22 kD, 27 kD and  $\gamma$  genes, could also be used. Other promoters known to function, for example, in maize include the promoters for the following genes: *waxy*, *Brittle*, *Shrunken 2*, Branching enzymes I and II, starch synthases, debranching enzymes, oleosins, glutelins and sucrose synthases. A particularly preferred promoter for maize endosperm expression is the promoter for the glutelin gene from rice, more particularly the Osgt-1 promoter (Zheng *et al.*, *Mol. Cell Biol.* 13:5829-5842 (1993), herein incorporated by reference in its entirety). Examples of promoters suitable for expression in wheat include those promoters for the ADPGlucose pyrosynthase (ADPGPP) subunits, the granule bound and other starch synthase, the branching and debranching enzymes, the embryogenesis-abundant proteins, the gliadins and the glutenins. Examples of such promoters in rice include those promoters for the ADPGPP subunits, the granule bound and other starch synthase, the branching enzymes, the debranching enzymes, sucrose synthases and the glutelins. A particularly preferred promoter is the promoter for rice glutelin, Osgt-1. Examples of such promoters for barley include those for the ADPGPP subunits, the granule bound and other starch synthase, the branching enzymes, the debranching enzymes, sucrose synthases, the hordeins, the embryo globulins and the aleurone specific proteins.

Root specific promoters may also be used. An example of such a promoter is the promoter for the acid chitinase gene (Samac *et al.*, *Plant Mol. Biol.* 25:587-596 (1994), the entirety of which is herein incorporated by reference). Expression in root tissue could also be accomplished by utilizing the root specific subdomains of the CaMV35S promoter that have been identified (Lam *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:7890-7894 (1989), herein incorporated by reference in its entirety). Other root cell specific promoters include those reported by Conkling *et al.* (Conkling *et al.*, *Plant Physiol.* 93:1203-1211 (1990), the entirety of which is herein incorporated by reference).

Additional promoters that may be utilized are described, for example, in U.S. Patent Nos. 5,378,619; 5,391,725; 5,428,147; 5,447,858; 5,608,144; 5,608,144; 5,614,399; 5,633,441; 5,633,435; and 4,633,436, all of which are herein incorporated in their entirety. In addition, a tissue specific enhancer may be used (Fromm *et al.*, *The Plant Cell* 1:977-984 (1989), the entirety of which is herein incorporated by reference).

Constructs or vectors may also include with the coding region of interest a nucleic acid sequence that acts, in whole or in part, to terminate transcription of that region. For example, such sequences have been isolated including the Tr7 3' sequence and the NOS 3' sequence (Ingelbrecht *et al.*, *The Plant Cell* 1:671-680 (1989), the entirety of which is herein incorporated by reference; Bevan *et al.*, *Nucleic Acids Res.* 11:369-385 (1983), the entirety of which is herein incorporated by reference), or the like.

A vector or construct may also include regulatory elements. Examples of such include the Adh intron 1 (Callis *et al.*, *Genes and Develop.* 1:1183-1200 (1987), the entirety of which is herein incorporated by reference), the sucrose synthase intron (Vasil *et al.*, *Plant Physiol.* 91:1575-1579 (1989), the entirety of which is herein incorporated by reference) and the TMV

omega element (Gallie *et al.*, *The Plant Cell* 1:301-311 (1989), the entirety of which is herein incorporated by reference). These and other regulatory elements may be included when appropriate.

A vector or construct may also include a selectable marker. Selectable markers may also be used to select for plants or plant cells that contain the exogenous genetic material. Examples of such include, but are not limited to, a neo gene (Potrykus *et al.*, *Mol. Gen. Genet.* 199:183-188 (1985), the entirety of which is herein incorporated by reference) which codes for kanamycin resistance and can be selected for using kanamycin, G418, etc.; a bar gene which codes for bialaphos resistance; a mutant EPSP synthase gene (Hinchey *et al.*, *Bio/Technology* 6:915-922 (1988), the entirety of which is herein incorporated by reference) which encodes glyphosate resistance; a nitrilase gene which confers resistance to bromoxynil (Stalker *et al.*, *J. Biol. Chem.* 263:6310-6314 (1988), the entirety of which is herein incorporated by reference); a mutant acetolactate synthase gene (ALS) which confers imidazolinone or sulphonylurea resistance (European Patent Application 154,204 (Sept. 11, 1985), the entirety of which is herein incorporated by reference); and a methotrexate resistant DHFR gene (Thillet *et al.*, *J. Biol. Chem.* 263:12500-12508 (1988), the entirety of which is herein incorporated by reference).

A vector or construct may also include a transit peptide. Incorporation of a suitable chloroplast transit peptide may also be employed (European Patent Application Publication Number 0218571, the entirety of which is herein incorporated by reference). Translational enhancers may also be incorporated as part of the vector DNA. DNA constructs could contain one or more 5' non-translated leader sequences which may serve to enhance expression of the gene products from the resulting mRNA transcripts. Such sequences may be derived from the promoter selected to express the gene or can be specifically modified to increase translation of

the mRNA. Such regions may also be obtained from viral RNAs, from suitable eukaryotic genes, or from a synthetic gene sequence. For a review of optimizing expression of transgenes, see Koziel *et al.*, *Plant Mol. Biol.* 32:393-405 (1996), the entirety of which is herein incorporated by reference.

A vector or construct may also include a screenable marker. Screenable markers may be used to monitor expression. Exemplary screenable markers include a  $\beta$ -glucuronidase or uidA gene (GUS) which encodes an enzyme for which various chromogenic substrates are known (Jefferson, *Plant Mol. Biol. Rep.* 5:387-405 (1987), the entirety of which is herein incorporated by reference; Jefferson *et al.*, *EMBO J.* 6:3901-3907 (1987), the entirety of which is herein incorporated by reference); an R-locus gene, which encodes a product that regulates the production of anthocyanin pigments (red color) in plant tissues (Dellaporta *et al.*, *Stadler Symposium* 11:263-282 (1988), the entirety of which is herein incorporated by reference); a  $\beta$ -lactamase gene (Sutcliffe *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 75:3737-3741 (1978), the entirety of which is herein incorporated by reference), a gene which encodes an enzyme for which various chromogenic substrates are known (e.g., PADAC, a chromogenic cephalosporin); a luciferase gene (Ow *et al.*, *Science* 234:856-859 (1986), the entirety of which is herein incorporated by reference); a xylE gene (Zukowsky *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 80:1101-1105 (1983), the entirety of which is herein incorporated by reference) which encodes a catechol dioxygenase that can convert chromogenic catechols; an  $\alpha$ -amylase gene (Ikata *et al.*, *Bio/Technol.* 8:241-242 (1990), the entirety of which is herein incorporated by reference); a tyrosinase gene (Katz *et al.*, *J. Gen. Microbiol.* 129:2703-2714 (1983), the entirety of which is herein incorporated by reference) which encodes an enzyme capable of oxidizing tyrosine to



DOPA and dopaquinone which in turn condenses to melanin; an  $\alpha$ -galactosidase, which will turn a chromogenic  $\alpha$ -galactose substrate.

Included within the terms “selectable or screenable marker genes” are also genes which encode a secretable marker whose secretion can be detected as a means of identifying or selecting for transformed cells. Examples include markers which encode a secretable antigen that can be identified by antibody interaction, or even secretable enzymes which can be detected catalytically. Secretable proteins fall into a number of classes, including small, diffusible proteins which are detectable, (*e.g.*, by ELISA), small active enzymes which are detectable in extracellular solution (*e.g.*,  $\alpha$ -amylase,  $\beta$ -lactamase, phosphinothricin transferase), or proteins which are inserted or trapped in the cell wall (such as proteins which include a leader sequence such as that found in the expression unit of extension or tobacco PR-S). Other possible selectable and/or screenable marker genes will be apparent to those of skill in the art.

There are many methods for introducing transforming nucleic acid molecules into plant cells. Suitable methods are believed to include virtually any method by which nucleic acid molecules may be introduced into a cell, such as by *Agrobacterium* infection or direct delivery of nucleic acid molecules such as, for example, by PEG-mediated transformation, by electroporation or by acceleration of DNA coated particles, etc (Potrykus, *Ann. Rev. Plant Physiol. Plant Mol. Biol.* 42:205-225 (1991), the entirety of which is herein incorporated by reference; Vasil, *Plant Mol. Biol.* 25:925-937 (1994), the entirety of which is herein incorporated by reference). For example, electroporation has been used to transform maize protoplasts (Fromm *et al.*, *Nature* 312:791-793 (1986), the entirety of which is herein incorporated by reference).

Other vector systems suitable for introducing transforming DNA into a host plant cell include but are not limited to binary artificial chromosome (BIBAC) vectors (Hamilton *et al.*, *Gene* 200:107-116 (1997), the entirety of which is herein incorporated by reference); and transfection with RNA viral vectors (Della-Cioppa *et al.*, *Ann. N.Y. Acad. Sci.* (1996), 792 (Engineering Plants for Commercial Products and Applications), 57-61, the entirety of which is herein incorporated by reference). Additional vector systems also include plant selectable YAC vectors such as those described in Mullen *et al.*, *Molecular Breeding* 4:449-457 (1988), the entirety of which is herein incorporated by reference).

Technology for introduction of DNA into cells is well known to those of skill in the art. Four general methods for delivering a gene into cells have been described: (1) chemical methods (Graham and van der Eb, *Virology* 54:536-539 (1973), the entirety of which is herein incorporated by reference); (2) physical methods such as microinjection (Capecchi, *Cell* 22:479-488 (1980), the entirety of which is herein incorporated by reference), electroporation (Wong and Neumann, *Biochem. Biophys. Res. Commun.* 107:584-587 (1982); Fromm *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 82:5824-5828 (1985); U.S. Patent No. 5,384,253, all of which are herein incorporated in their entirety); and the gene gun (Johnston and Tang, *Methods Cell Biol.* 43:353-365 (1994), the entirety of which is herein incorporated by reference); (3) viral vectors (Clapp, *Clin. Perinatol.* 20:155-168 (1993); Lu *et al.*, *J. Exp. Med.* 178:2089-2096 (1993); Eglitis and Anderson, *Biotechniques* 6:608-614 (1988), all of which are herein incorporated in their entirety); and (4) receptor-mediated mechanisms (Curiel *et al.*, *Hum. Gen. Ther.* 3:147-154 (1992), Wagner *et al.*, *Proc. Natl. Acad. Sci. (USA)* 89:6099-6103 (1992), both of which are incorporated by reference in their entirety).

Acceleration methods that may be used include, for example, microprojectile bombardment and the like. One example of a method for delivering transforming nucleic acid molecules to plant cells is microprojectile bombardment. This method has been reviewed by Yang and Christou (eds.), *Particle Bombardment Technology for Gene Transfer*, Oxford Press, Oxford, England (1994), the entirety of which is herein incorporated by reference). Non-biological particles (microprojectiles) that may be coated with nucleic acids and delivered into cells by a propelling force. Exemplary particles include those comprised of tungsten, gold, platinum and the like.

A particular advantage of microprojectile bombardment, in addition to it being an effective means of reproducibly transforming monocots, is that neither the isolation of protoplasts (Cristou *et al.*, *Plant Physiol.* 87:671-674 (1988), the entirety of which is herein incorporated by reference) nor the susceptibility of *Agrobacterium* infection are required. An illustrative embodiment of a method for delivering DNA into maize cells by acceleration is a biolistics  $\alpha$ -particle delivery system, which can be used to propel particles coated with DNA through a screen, such as a stainless steel or Nytex screen, onto a filter surface covered with corn cells cultured in suspension. Gordon-Kamm *et al.*, describes the basic procedure for coating tungsten particles with DNA (Gordon-Kamm *et al.*, *Plant Cell* 2:603-618 (1990), the entirety of which is herein incorporated by reference). The screen disperses the tungsten nucleic acid particles so that they are not delivered to the recipient cells in large aggregates. A particle delivery system suitable for use with the present invention is the helium acceleration PDS-1000/He gun is available from Bio-Rad Laboratories (Bio-Rad, Hercules, California)(Sanford *et al.*, *Technique* 3:3-16 (1991), the entirety of which is herein incorporated by reference).

For the bombardment, cells in suspension may be concentrated on filters. Filters containing the cells to be bombarded are positioned at an appropriate distance below the microprojectile stopping plate. If desired, one or more screens are also positioned between the gun and the cells to be bombarded.

Alternatively, immature embryos or other target cells may be arranged on solid culture medium. The cells to be bombarded are positioned at an appropriate distance below the microprojectile stopping plate. If desired, one or more screens are also positioned between the acceleration device and the cells to be bombarded. Through the use of techniques set forth herein one may obtain up to 1000 or more foci of cells transiently expressing a marker gene. The number of cells in a focus which express the exogenous gene product 48 hours post-bombardment often range from one to ten and average one to three.

In bombardment transformation, one may optimize the pre-bombardment culturing conditions and the bombardment parameters to yield the maximum numbers of stable transformants. Both the physical and biological parameters for bombardment are important in this technology. Physical factors are those that involve manipulating the DNA/microprojectile precipitate or those that affect the flight and velocity of either the macro- or microprojectiles. Biological factors include all steps involved in manipulation of cells before and immediately after bombardment, the osmotic adjustment of target cells to help alleviate the trauma associated with bombardment and also the nature of the transforming DNA, such as linearized DNA or intact supercoiled plasmids. It is believed that pre-bombardment manipulations are especially important for successful transformation of immature embryos.

In another alternative embodiment, plastids can be stably transformed. Methods disclosed for plastid transformation in higher plants include the particle gun delivery of DNA

containing a selectable marker and targeting of the DNA to the plastid genome through homologous recombination (Svab *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 87:8526-8530 (1990); Svab and Maliga, *Proc. Natl. Acad. Sci. (U.S.A.)* 90:913-917 (1993); Staub and Maliga, *EMBO J.* 12:601-606 (1993); U.S. Patents 5, 451,513 and 5,545,818, all of which are herein incorporated by reference in their entirety).

Accordingly, it is contemplated that one may wish to adjust various aspects of the bombardment parameters in small scale studies to fully optimize the conditions. One may particularly wish to adjust physical parameters such as gap distance, flight distance, tissue distance and helium pressure. One may also minimize the trauma reduction factors by modifying conditions which influence the physiological state of the recipient cells and which may therefore influence transformation and integration efficiencies. For example, the osmotic state, tissue hydration and the subculture stage or cell cycle of the recipient cells may be adjusted for optimum transformation. The execution of other routine adjustments will be known to those of skill in the art in light of the present disclosure.

*Agrobacterium*-mediated transfer is a widely applicable system for introducing genes into plant cells because the DNA can be introduced into whole plant tissues, thereby bypassing the need for regeneration of an intact plant from a protoplast. The use of *Agrobacterium*-mediated plant integrating vectors to introduce DNA into plant cells is well known in the art. See, for example the methods described by Fraley *et al.*, *Bio/Technology* 3:629-635 (1985) and Rogers *et al.*, *Methods Enzymol.* 153:253-277 (1987), both of which are herein incorporated by reference in their entirety. Further, the integration of the Ti-DNA is a relatively precise process resulting in few rearrangements. The region of DNA to be transferred is defined by the border sequences and

intervening DNA is usually inserted into the plant genome as described (Spielmann *et al.*, *Mol. Gen. Genet.* 205:34 (1986), the entirety of which is herein incorporated by reference).

Modern *Agrobacterium* transformation vectors are capable of replication in *E. coli* as well as *Agrobacterium*, allowing for convenient manipulations as described (Klee *et al.*, In: *Plant DNA Infectious Agents*, Hohn and Schell (eds.), Springer-Verlag, New York, pp. 179-203 (1985), the entirety of which is herein incorporated by reference. Moreover, technological advances in vectors for *Agrobacterium*-mediated gene transfer have improved the arrangement of genes and restriction sites in the vectors to facilitate construction of vectors capable of expressing various polypeptide coding genes. The vectors described have convenient multi-linker regions flanked by a promoter and a polyadenylation site for direct expression of inserted polypeptide coding genes and are suitable for present purposes (Rogers *et al.*, *Methods Enzymol.* 153:253-277 (1987)). In addition, *Agrobacterium* containing both armed and disarmed Ti genes can be used for the transformations. In those plant strains where *Agrobacterium*-mediated transformation is efficient, it is the method of choice because of the facile and defined nature of the gene transfer.

A transgenic plant formed using *Agrobacterium* transformation methods typically contains a single gene on one chromosome. Such transgenic plants can be referred to as being heterozygous for the added gene. More preferred is a transgenic plant that is homozygous for the added structural gene; *i.e.*, a transgenic plant that contains two added genes, one gene at the same locus on each chromosome of a chromosome pair. A homozygous transgenic plant can be obtained by sexually mating (selfing) an independent segregant transgenic plant that contains a single added gene, germinating some of the seed produced and analyzing the resulting plants produced for the gene of interest.

It is also to be understood that two different transgenic plants can also be mated to produce offspring that contain two independently segregating added, exogenous genes. Selfing of appropriate progeny can produce plants that are homozygous for both added, exogenous genes that encode a polypeptide of interest. Back-crossing to a parental plant and out-crossing with a non-transgenic plant are also contemplated, as is vegetative propagation.

Transformation of plant protoplasts can be achieved using methods based on calcium phosphate precipitation, polyethylene glycol treatment, electroporation and combinations of these treatments (*See, for example*, Potrykus *et al.*, *Mol. Gen. Genet.* 205:193-200 (1986); Lorz *et al.*, *Mol. Gen. Genet.* 199:178 (1985); Fromm *et al.*, *Nature* 319:791 (1986); Uchimiya *et al.*, *Mol. Gen. Genet.* 204:204 (1986); Marcotte *et al.*, *Nature* 335:454-457 (1988), all of which are herein incorporated by reference in their entirety).

Application of these systems to different plant strains depends upon the ability to regenerate that particular plant strain from protoplasts. Illustrative methods for the regeneration of cereals from protoplasts are described (Fujimura *et al.*, *Plant Tissue Culture Letters* 2:74 (1985); Toriyama *et al.*, *Theor Appl. Genet.* 205:34 (1986); Yamada *et al.*, *Plant Cell Rep.* 4:85 (1986); Abdullah *et al.*, *Biotechnolog* 4:1087 (1986), all of which are herein incorporated by reference in their entirety).

To transform plant strains that cannot be successfully regenerated from protoplasts, other ways to introduce DNA into intact cells or tissues can be utilized. For example, regeneration of cereals from immature embryos or explants can be effected as described (Vasil, *Biotechnology* 6:397 (1988), the entirety of which is herein incorporated by reference). In addition, "particle gun" or high-velocity microprojectile technology can be utilized (Vasil *et al.*, *Bio/Technology* 10:667 (1992), the entirety of which is herein incorporated by reference).

Using the latter technology, DNA is carried through the cell wall and into the cytoplasm on the surface of small metal particles as described (Klein *et al.*, *Nature* 328:70 (1987); Klein *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 85:8502-8505 (1988); McCabe *et al.*, *Bio/Technology* 6:923 (1988), all of which are herein incorporated by reference in their entirety). The metal particles penetrate through several layers of cells and thus allow the transformation of cells within tissue explants.

Other methods of cell transformation can also be used and include but are not limited to introduction of DNA into plants by direct DNA transfer into pollen (Zhou *et al.*, *Methods Enzymol.* 101:433 (1983); Hess *et al.*, *Intern Rev. Cytol.* 107:367 (1987); Luo *et al.*, *Plant Mol Biol. Reporter* 6:165 (1988), all of which are herein incorporated by reference in their entirety), by direct injection of DNA into reproductive organs of a plant (Pena *et al.*, *Nature* 325:274 (1987), the entirety of which is herein incorporated by reference), or by direct injection of DNA into the cells of immature embryos followed by the rehydration of desiccated embryos (Neuhaus *et al.*, *Theor. Appl. Genet.* 75:30 (1987), the entirety of which is herein incorporated by reference).

The regeneration, development and cultivation of plants from single plant protoplast transformants or from various transformed explants is well known in the art (Weissbach and Weissbach, In: *Methods for Plant Molecular Biology*, Academic Press, San Diego, CA, (1988), the entirety of which is herein incorporated by reference). This regeneration and growth process typically includes the steps of selection of transformed cells, culturing those individualized cells through the usual stages of embryonic development through the rooted plantlet stage. Transgenic embryos and seeds are similarly regenerated. The resulting transgenic rooted shoots are thereafter planted in an appropriate plant growth medium such as soil.



The development or regeneration of plants containing the foreign, exogenous gene that encodes a protein of interest is well known in the art. Preferably, the regenerated plants are self-pollinated to provide homozygous transgenic plants. Otherwise, pollen obtained from the regenerated plants is crossed to seed-grown plants of agronomically important lines. Conversely, pollen from plants of these important lines is used to pollinate regenerated plants. A transgenic plant of the present invention containing a desired polypeptide is cultivated using methods well known to one skilled in the art.

There are a variety of methods for the regeneration of plants from plant tissue. The particular method of regeneration will depend on the starting plant tissue and the particular plant species to be regenerated.

Methods for transforming dicots, primarily by use of *Agrobacterium tumefaciens* and obtaining transgenic plants have been published for cotton (U.S. Patent No. 5,004,863; U.S. Patent No. 5,159,135; U.S. Patent No. 5,518,908, all of which are herein incorporated by reference in their entirety); soybean (U.S. Patent No. 5,569,834; U.S. Patent No. 5,416,011; McCabe *et al.*, *Biotechnology* 6:923 (1988); Christou *et al.*, *Plant Physiol.* 87:671-674 (1988); all of which are herein incorporated by reference in their entirety); *Brassica* (U.S. Patent No. 5,463,174, the entirety of which is herein incorporated by reference); peanut (Cheng *et al.*, *Plant Cell Rep.* 15:653-657 (1996), McKently *et al.*, *Plant Cell Rep.* 14:699-703 (1995), all of which are herein incorporated by reference in their entirety); papaya; and pea (Grant *et al.*, *Plant Cell Rep.* 15:254-258 (1995), the entirety of which is herein incorporated by reference).

Transformation of monocotyledons using electroporation, particle bombardment and *Agrobacterium* have also been reported. Transformation and plant regeneration have been achieved in asparagus (Bytebier *et al.*, *Proc. Natl. Acad. Sci. (USA)* 84:5354 (1987), the entirety

of which is herein incorporated by reference); barley (Wan and Lemaux, *Plant Physiol* 104:37 (1994), the entirety of which is herein incorporated by reference); maize (Rhodes *et al.*, *Science* 240:204 (1988); Gordon-Kamm *et al.*, *Plant Cell* 2:603-618 (1990); Fromm *et al.*, *Bio/Technology* 8:833 (1990); Koziel *et al.*, *Bio/Technology* 11:194 (1993); Armstrong *et al.*, *Crop Science* 35:550-557 (1995); all of which are herein incorporated by reference in their entirety); oat (Somers *et al.*, *Bio/Technology* 10:1589 (1992), the entirety of which is herein incorporated by reference); orchard grass (Horn *et al.*, *Plant Cell Rep.* 7:469 (1988), the entirety of which is herein incorporated by reference); rice (Toriyama *et al.*, *Theor Appl. Genet.* 205:34 (1986); Part *et al.*, *Plant Mol. Biol.* 32:1135-1148 (1996); Abedinia *et al.*, *Aust. J. Plant Physiol.* 24:133-141 (1997); Zhang and Wu, *Theor. Appl. Genet.* 76:835 (1988); Zhang *et al.*, *Plant Cell Rep.* 7:379 (1988); Battra and Hall, *Plant Sci.* 86:191-202 (1992); Christou *et al.*, *Bio/Technology* 9:957 (1991), all of which are herein incorporated by reference in their entirety); rye (De la Pena *et al.*, *Nature* 325:274 (1987), the entirety of which is herein incorporated by reference); sugarcane (Bower and Birch, *Plant J.* 2:409 (1992), the entirety of which is herein incorporated by reference); tall fescue (Wang *et al.*, *Bio/Technology* 10:691 (1992), the entirety of which is herein incorporated by reference) and wheat (Vasil *et al.*, *Bio/Technology* 10:667 (1992), the entirety of which is herein incorporated by reference; U.S. Patent No. 5,631,152, the entirety of which is herein incorporated by reference.)

Assays for gene expression based on the transient expression of cloned nucleic acid constructs have been developed by introducing the nucleic acid molecules into plant cells by polyethylene glycol treatment, electroporation, or particle bombardment (Marcotte *et al.*, *Nature* 335:454-457 (1988), the entirety of which is herein incorporated by reference; Marcotte *et al.*, *Plant Cell* 1:523-532 (1989), the entirety of which is herein incorporated by reference; McCarty

*et al.*, *Cell* 66:895-905 (1991), the entirety of which is herein incorporated by reference; Hattori *et al.*, *Genes Dev.* 6:609-618 (1992), the entirety of which is herein incorporated by reference; Goff *et al.*, *EMBO J.* 9:2517-2522 (1990), the entirety of which is herein incorporated by reference). Transient expression systems may be used to functionally dissect gene constructs (see generally, Mailga *et al.*, *Methods in Plant Molecular Biology*, Cold Spring Harbor Press (1995)).

Any of the nucleic acid molecules of the present invention may be introduced into a plant cell in a permanent or transient manner in combination with other genetic elements such as vectors, promoters, enhancers etc. Further, any of the nucleic acid molecules of the present invention may be introduced into a plant cell in a manner that allows for overexpression of the protein or fragment thereof encoded by the nucleic acid molecule.

Cosuppression is the reduction in expression levels, usually at the level of RNA, of a particular endogenous gene or gene family by the expression of a homologous sense construct that is capable of transcribing mRNA of the same strandedness as the transcript of the endogenous gene (Napoli *et al.*, *Plant Cell* 2:279-289 (1990), the entirety of which is herein incorporated by reference; van der Krol *et al.*, *Plant Cell* 2:291-299 (1990), the entirety of which is herein incorporated by reference). Cosuppression may result from stable transformation with a single copy nucleic acid molecule that is homologous to a nucleic acid sequence found with the cell (Prolls and Meyer, *Plant J.* 2:465-475 (1992), the entirety of which is herein incorporated by reference) or with multiple copies of a nucleic acid molecule that is homologous to a nucleic acid sequence found with the cell (Mittlesten *et al.*, *Mol. Gen. Genet.* 244:325-330 (1994), the entirety of which is herein incorporated by reference). Genes, even though different, linked to

homologous promoters may result in the cosuppression of the linked genes (Vaucheret, *C.R. Acad. Sci. III* 316:1471-1483 (1993), the entirety of which is herein incorporated by reference).

This technique has, for example, been applied to generate white flowers from red petunia and tomatoes that do not ripen on the vine. Up to 50% of petunia transformants that contained a sense copy of the glucoamylase (CHS) gene produced white flowers or floral sectors; this was as a result of the post-transcriptional loss of mRNA encoding CHS (Flavell, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:3490-3496 (1994), the entirety of which is herein incorporated by reference); van Blokland *et al.*, *Plant J.* 6:861-877 (1994), the entirety of which is herein incorporated by reference). Cosuppression may require the coordinate transcription of the transgene and the endogenous gene and can be reset by a developmental control mechanism (Jorgensen, *Trends Biotechnol.* 8:340-344 (1990), the entirety of which is herein incorporated by reference; Meins and Kunz, In: *Gene Inactivation and Homologous Recombination in Plants*, Paszkowski (ed.), pp. 335-348, Kluwer Academic, Netherlands (1994), the entirety of which is herein incorporated by reference).

It is understood that one or more of the nucleic acids of the present invention may be introduced into a plant cell and transcribed using an appropriate promoter with such transcription resulting in the cosuppression of an endogenous methionine pathway protein.

Antisense approaches are a way of preventing or reducing gene function by targeting the genetic material (Mol *et al.*, *FEBS Lett.* 268:427-430 (1990), the entirety of which is herein incorporated by reference). The objective of the antisense approach is to use a sequence complementary to the target gene to block its expression and create a mutant cell line or organism in which the level of a single chosen protein is selectively reduced or abolished. Antisense techniques have several advantages over other 'reverse genetic' approaches. The site

of inactivation and its developmental effect can be manipulated by the choice of promoter for antisense genes or by the timing of external application or microinjection. Antisense can manipulate its specificity by selecting either unique regions of the target gene or regions where it shares homology to other related genes (Hiatt *et al.*, In: *Genetic Engineering*, Setlow (ed.), Vol. 11, New York: Plenum 49-63 (1989), the entirety of which is herein incorporated by reference).

The principle of regulation by antisense RNA is that RNA that is complementary to the target mRNA is introduced into cells, resulting in specific RNA:RNA duplexes being formed by base pairing between the antisense substrate and the target mRNA (Green *et al.*, *Annu. Rev. Biochem.* 55:569-597 (1986), the entirety of which is herein incorporated by reference). Under one embodiment, the process involves the introduction and expression of an antisense gene sequence. Such a sequence is one in which part or all of the normal gene sequences are placed under a promoter in inverted orientation so that the 'wrong' or complementary strand is transcribed into a noncoding antisense RNA that hybridizes with the target mRNA and interferes with its expression (Takayama and Inouye, *Crit. Rev. Biochem. Mol. Biol.* 25:155-184 (1990), the entirety of which is herein incorporated by reference). An antisense vector is constructed by standard procedures and introduced into cells by transformation, transfection, electroporation, microinjection, infection, etc. The type of transformation and choice of vector will determine whether expression is transient or stable. The promoter used for the antisense gene may influence the level, timing, tissue, specificity, or inducibility of the antisense inhibition.

It is understood that the activity of a methionine pathway protein in a plant cell may be reduced or depressed by growing a transformed plant cell containing a nucleic acid molecule whose non-transcribed strand encodes a methionine pathway protein or fragment thereof.

Antibodies have been expressed in plants (Hiatt *et al.*, *Nature* 342:76-78 (1989), the entirety of which is herein incorporated by reference; Conrad and Fielder, *Plant Mol. Biol.* 26:1023-1030 (1994), the entirety of which is herein incorporated by reference). Cytoplasmic expression of a scFv (single-chain Fv antibodies) has been reported to delay infection by artichoke mottled crinkle virus. Transgenic plants that express antibodies directed against endogenous proteins may exhibit a physiological effect (Philips *et al.*, *EMBO J.* 16:4489-4496 (1997), the entirety of which is herein incorporated by reference; Marion-Poll, *Trends in Plant Science* 2:447-448 (1997), the entirety of which is herein incorporated by reference). For example, expressed anti-abscisic antibodies have been reported to result in a general perturbation of seed development (Philips *et al.*, *EMBO J.* 16: 4489-4496 (1997)).

Antibodies that are catalytic may also be expressed in plants (abzymes). The principle behind abzymes is that since antibodies may be raised against many molecules, this recognition ability can be directed toward generating antibodies that bind transition states to force a chemical reaction forward (Persidas, *Nature Biotechnology* 15:1313-1315 (1997), the entirety of which is herein incorporated by reference; Baca *et al.*, *Ann. Rev. Biophys. Biomol. Struct.* 26:461-493 (1997), the entirety of which is herein incorporated by reference). The catalytic abilities of abzymes may be enhanced by site directed mutagenesis. Examples of abzymes are, for example, set forth in U.S. Patent No: 5,658,753; U.S. Patent No. 5,632,990; U.S. Patent No. 5,631,137; U.S. Patent 5,602,015; U.S. Patent No. 5,559,538; U.S. Patent No. 5,576,174; U.S. Patent No. 5,500,358; U.S. Patent 5,318,897; U.S. Patent No. 5,298,409; U.S. Patent No. 5,258,289 and U.S. Patent No. 5,194,585, all of which are herein incorporated in their entirety.

It is understood that any of the antibodies of the present invention may be expressed in plants and that such expression can result in a physiological effect. It is also understood that any of the expressed antibodies may be catalytic.

**(b) Fungal Constructs and Fungal Transformants**

The present invention also relates to a fungal recombinant vector comprising exogenous genetic material. The present invention also relates to a fungal cell comprising a fungal recombinant vector. The present invention also relates to methods for obtaining a recombinant fungal host cell comprising introducing into a fungal host cell exogenous genetic material.

Exogenous genetic material may be transferred into a fungal cell. In a preferred embodiment the exogenous genetic material includes a nucleic acid molecule of the present invention having a sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof or fragments of either. The fungal recombinant vector may be any vector which can be conveniently subjected to recombinant DNA procedures. The choice of a vector will typically depend on the compatibility of the vector with the fungal host cell into which the vector is to be introduced. The vector may be a linear or a closed circular plasmid. The vector system may be a single vector or plasmid or two or more vectors or plasmids which together contain the total DNA to be introduced into the genome of the fungal host.

The fungal vector may be an autonomously replicating vector, *i.e.*, a vector which exists as an extrachromosomal entity, the replication of which is independent of chromosomal replication, *e.g.*, a plasmid, an extrachromosomal element, a minichromosome, or an artificial chromosome. The vector may contain any means for assuring self-replication. Alternatively, the vector may be one which, when introduced into the fungal cell, is integrated into the genome and replicated together with the chromosome(s) into which it has been integrated. For integration,

the vector may rely on the nucleic acid sequence of the vector for stable integration of the vector into the genome by homologous or nonhomologous recombination. Alternatively, the vector may contain additional nucleic acid sequences for directing integration by homologous recombination into the genome of the fungal host. The additional nucleic acid sequences enable the vector to be integrated into the host cell genome at a precise location(s) in the chromosome(s). To increase the likelihood of integration at a precise location, there should be preferably two nucleic acid sequences which individually contain a sufficient number of nucleic acids, preferably 400bp to 1500bp, more preferably 800bp to 1000bp, which are highly homologous with the corresponding target sequence to enhance the probability of homologous recombination. These nucleic acid sequences may be any sequence that is homologous with a target sequence in the genome of the fungal host cell and, furthermore, may be non-encoding or encoding sequences.

For autonomous replication, the vector may further comprise an origin of replication enabling the vector to replicate autonomously in the host cell in question. Examples of origin of replications for use in a yeast host cell are the 2 micron origin of replication and the combination of CEN3 and ARS 1. Any origin of replication may be used which is compatible with the fungal host cell of choice.

The fungal vectors of the present invention preferably contain one or more selectable markers which permit easy selection of transformed cells. A selectable marker is a gene the product of which provides, for example biocide or viral resistance, resistance to heavy metals, prototrophy to auxotrophs and the like. The selectable marker may be selected from the group including, but not limited to, *amdS* (acetamidase), *argB* (ornithine carbamoyltransferase), *bar* (phosphinothricin acetyltransferase), *hygB* (hygromycin phosphotransferase), *niaD* (nitrate



reductase), *pyrG* (orotidine-5'-phosphate decarboxylase) and *sC* (sulfate adenylyltransferase) and *trpC* (anthranilate synthase). Preferred for use in an *Aspergillus* cell are the *amdS* and *pyrG* markers of *Aspergillus nidulans* or *Aspergillus oryzae* and the bar marker of *Streptomyces hygroscopicus*. Furthermore, selection may be accomplished by co-transformation, *e.g.*, as described in WO 91/17243, the entirety of which is herein incorporated by reference. A nucleic acid sequence of the present invention may be operably linked to a suitable promoter sequence. The promoter sequence is a nucleic acid sequence which is recognized by the fungal host cell for expression of the nucleic acid sequence. The promoter sequence contains transcription and translation control sequences which mediate the expression of the protein or fragment thereof.

A promoter may be any nucleic acid sequence which shows transcriptional activity in the fungal host cell of choice and may be obtained from genes encoding polypeptides either homologous or heterologous to the host cell. Examples of suitable promoters for directing the transcription of a nucleic acid construct of the invention in a filamentous fungal host are promoters obtained from the genes encoding *Aspergillus oryzae* TAKA amylase, *Rhizomucor miehei* aspartic proteinase, *Aspergillus niger* neutral alpha-amylase, *Aspergillus niger* acid stable alpha-amylase, *Aspergillus niger* or *Aspergillus awamori* glucoamylase (*glaA*), *Rhizomucor miehei* lipase, *Aspergillus oryzae* alkaline protease, *Aspergillus oryzae* triose phosphate isomerase, *Aspergillus nidulans* acetamidase and hybrids thereof. In a yeast host, a useful promoter is the *Saccharomyces cerevisiae* enolase (*eno-1*) promoter. Particularly preferred promoters are the TAKA amylase, NA2-tpi (a hybrid of the promoters from the genes encoding *Aspergillus niger* neutral alpha -amylase and *Aspergillus oryzae* triose phosphate isomerase) and *glaA* promoters.

A protein or fragment thereof encoding nucleic acid molecule of the present invention may also be operably linked to a terminator sequence at its 3' terminus. The terminator sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any terminator which is functional in the fungal host cell of choice may be used in the present invention, but particularly preferred terminators are obtained from the genes encoding *Aspergillus oryzae* TAKA amylase, *Aspergillus niger* glucoamylase, *Aspergillus nidulans* anthranilate synthase, *Aspergillus niger* alpha-glucosidase and *Saccharomyces cerevisiae* enolase.

A protein or fragment thereof encoding nucleic acid molecule of the present invention may also be operably linked to a suitable leader sequence. A leader sequence is a nontranslated region of a mRNA which is important for translation by the fungal host. The leader sequence is operably linked to the 5' terminus of the nucleic acid sequence encoding the protein or fragment thereof. The leader sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any leader sequence which is functional in the fungal host cell of choice may be used in the present invention, but particularly preferred leaders are obtained from the genes encoding *Aspergillus oryzae* TAKA amylase and *Aspergillus oryzae* triose phosphate isomerase.

A polyadenylation sequence may also be operably linked to the 3' terminus of the nucleic acid sequence of the present invention. The polyadenylation sequence is a sequence which when transcribed is recognized by the fungal host to add polyadenosine residues to transcribed mRNA. The polyadenylation sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any polyadenylation sequence which is functional in the fungal host of choice may be used in the present invention, but particularly

preferred polyadenylation sequences are obtained from the genes encoding *Aspergillus oryzae* TKA amylase, *Aspergillus niger* glucoamylase, *Aspergillus nidulans* anthranilate synthase and *Aspergillus niger* alpha-glucosidase.

To avoid the necessity of disrupting the cell to obtain the protein or fragment thereof and to minimize the amount of possible degradation of the expressed protein or fragment thereof within the cell, it is preferred that expression of the protein or fragment thereof gives rise to a product secreted outside the cell. To this end, a protein or fragment thereof of the present invention may be linked to a signal peptide linked to the amino terminus of the protein or fragment thereof. A signal peptide is an amino acid sequence which permits the secretion of the protein or fragment thereof from the fungal host into the culture medium. The signal peptide may be native to the protein or fragment thereof of the invention or may be obtained from foreign sources. The 5' end of the coding sequence of the nucleic acid sequence of the present invention may inherently contain a signal peptide coding region naturally linked in translation reading frame with the segment of the coding region which encodes the secreted protein or fragment thereof. Alternatively, the 5' end of the coding sequence may contain a signal peptide coding region which is foreign to that portion of the coding sequence which encodes the secreted protein or fragment thereof. The foreign signal peptide may be required where the coding sequence does not normally contain a signal peptide coding region. Alternatively, the foreign signal peptide may simply replace the natural signal peptide to obtain enhanced secretion of the desired protein or fragment thereof. The foreign signal peptide coding region may be obtained from a glucoamylase or an amylase gene from an *Aspergillus* species, a lipase or proteinase gene from *Rhizomucor miehei*, the gene for the alpha-factor from *Saccharomyces cerevisiae*, or the calf preprochymosin gene. An effective signal peptide for fungal host cells is the *Aspergillus*

*oryzae* TAKA amylase signal, *Aspergillus niger* neutral amylase signal, the *Rhizomucor miehei* aspartic proteinase signal, the *Humicola lanuginosus* cellulase signal, or the *Rhizomucor miehei* lipase signal. However, any signal peptide capable of permitting secretion of the protein or fragment thereof in a fungal host of choice may be used in the present invention.

A protein or fragment thereof encoding nucleic acid molecule of the present invention may also be linked to a propeptide coding region. A propeptide is an amino acid sequence found at the amino terminus of a protein or proenzyme. Cleavage of the propeptide from the proprotein yields a mature biochemically active protein. The resulting polypeptide is known as a propeptide or proenzyme (or a zymogen in some cases). Propeptides are generally inactive and can be converted to mature active polypeptides by catalytic or autocatalytic cleavage of the propeptide from the propeptide or proenzyme. The propeptide coding region may be native to the protein or fragment thereof or may be obtained from foreign sources. The foreign propeptide coding region may be obtained from the *Saccharomyces cerevisiae* alpha-factor gene or *Myceliophthora thermophila* laccase gene (WO 95/33836, the entirety of which is herein incorporated by reference).

The procedures used to ligate the elements described above to construct the recombinant expression vector of the present invention are well known to one skilled in the art (see, for example, Sambrook *et al.*, *Molecular Cloning, A Laboratory Manual*, 2nd ed., Cold Spring Harbor, N.Y., (1989)).

The present invention also relates to recombinant fungal host cells produced by the methods of the present invention which are advantageously used with the recombinant vector of the present invention. The cell is preferably transformed with a vector comprising a nucleic acid sequence of the invention followed by integration of the vector into the host chromosome. The

choice of fungal host cells will to a large extent depend upon the gene encoding the protein or fragment thereof and its source. The fungal host cell may, for example, be a yeast cell or a filamentous fungal cell.

"Yeast" as used herein includes *Ascosporogenous* yeast (*Endomycetales*), *Basidiosporogenous* yeast and yeast belonging to the *Fungi Imperfecti* (*Blastomycetes*). The *Ascosporogenous* yeasts are divided into the families *Spermophthoraceae* and *Saccharomycetaceae*. The latter is comprised of four subfamilies, *Schizosaccharomycoideae* (for example, genus *Schizosaccharomyces*), *Nadsonioideae*, *Lipomycoideae* and *Saccharomycoideae* (for example, genera *Pichia*, *Kluyveromyces* and *Saccharomyces*). The *Basidiosporogenous* yeasts include the genera *Leucosporidim*, *Rhodospordium*, *Sporidiobolus*, *Filobasidium* and *Filobasidiella*. Yeast belonging to the *Fungi Imperfecti* are divided into two families, *Sporobolomycetaceae* (for example, genera *Sorobolomyces* and *Bullera*) and *Cryptococcaceae* (for example, genus *Candida*). Since the classification of yeast may change in the future, for the purposes of this invention, yeast shall be defined as described in Biology and Activities of Yeast (Skinner *et al.*, *Soc. App. Bacteriol. Symposium Series* No. 9, (1980), the entirety of which is herein incorporated by reference). The biology of yeast and manipulation of yeast genetics are well known in the art (*see*, for example, *Biochemistry and Genetics of Yeast*, Bacil *et al.* (ed.), 2nd edition, 1987; *The Yeasts*, Rose and Harrison (eds.), 2nd ed., (1987); and *The Molecular Biology of the Yeast Saccharomyces*, Strathern *et al.* (eds.), (1981), all of which are herein incorporated by reference in their entirety).

"Fungi" as used herein includes the phyla *Ascomycota*, *Basidiomycota*, *Chytridiomycota* and *Zygomycota* (as defined by Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK; the entirety of

which is herein incorporated by reference) as well as the *Oomycota* (as cited in Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK) and all mitosporic fungi (Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK). Representative groups of *Ascomycota* include, for example, *Neurospora*, *Eupenicillium* (= *Penicillium*), *Emericella* (= *Aspergillus*), *Eurotium* (= *Aspergillus*) and the true yeasts listed above. Examples of *Basidiomycota* include mushrooms, rusts and smuts.

Representative groups of *Chytridiomycota* include, for example, *Allomyces*, *Blastocladiella*, *Coelomomyces* and aquatic fungi. Representative groups of *Oomycota* include, for example, *Saprolegniomycetous* aquatic fungi (water molds) such as *Achlya*. Examples of mitosporic fungi include *Aspergillus*, *Penicillium*, *Candida* and *Alternaria*. Representative groups of *Zygomycota* include, for example, *Rhizopus* and *Mucor*.

"Filamentous fungi" include all filamentous forms of the subdivision *Eumycota* and *Oomycota* (as defined by Hawksworth *et al.*, In: Ainsworth and Bisby's *Dictionary of The Fungi*, 8th edition, 1995, CAB International, University Press, Cambridge, UK). The filamentous fungi are characterized by a vegetative mycelium composed of chitin, cellulose, glucan, chitosan, mannan and other complex polysaccharides. Vegetative growth is by hyphal elongation and carbon catabolism is obligately aerobic. In contrast, vegetative growth by yeasts such as *Saccharomyces cerevisiae* is by budding of a unicellular thallus and carbon catabolism may be fermentative.

In one embodiment, the fungal host cell is a yeast cell. In a preferred embodiment, the yeast host cell is a cell of the species of *Candida*, *Kluyveromyces*, *Saccharomyces*, *Schizosaccharomyces*, *Pichia* and *Yarrowia*. In a preferred embodiment, the yeast host cell is a

*Saccharomyces cerevisiae* cell, a *Saccharomyces carlsbergensis*, *Saccharomyces diastaticus* cell, a *Saccharomyces douglasii* cell, a *Saccharomyces kluyveri* cell, a *Saccharomyces norbensis* cell, or a *Saccharomyces oviformis* cell. In another preferred embodiment, the yeast host cell is a *Kluyveromyces lactis* cell. In another preferred embodiment, the yeast host cell is a *Yarrowia lipolytica* cell.

In another embodiment, the fungal host cell is a filamentous fungal cell. In a preferred embodiment, the filamentous fungal host cell is a cell of the species of, but not limited to, *Acremonium*, *Aspergillus*, *Fusarium*, *Humicola*, *Myceliophthora*, *Mucor*, *Neurospora*, *Penicillium*, *Thielavia*, *Tolypocladium* and *Trichoderma*. In a preferred embodiment, the filamentous fungal host cell is an *Aspergillus* cell. In another preferred embodiment, the filamentous fungal host cell is an *Acremonium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Fusarium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Humicola* cell. In another preferred embodiment, the filamentous fungal host cell is a *Myceliophthora* cell. In another even preferred embodiment, the filamentous fungal host cell is a *Mucor* cell. In another preferred embodiment, the filamentous fungal host cell is a *Neurospora* cell. In another preferred embodiment, the filamentous fungal host cell is a *Penicillium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Thielavia* cell. In another preferred embodiment, the filamentous fungal host cell is a *Tolypocladium* cell. In another preferred embodiment, the filamentous fungal host cell is a *Trichoderma* cell. In a preferred embodiment, the filamentous fungal host cell is an *Aspergillus oryzae* cell, an *Aspergillus niger* cell, an *Aspergillus foetidus* cell, or an *Aspergillus japonicus* cell. In another preferred embodiment, the filamentous fungal host cell is a *Fusarium oxysporum* cell or a *Fusarium graminearum* cell. In another preferred embodiment, the filamentous fungal

host cell is a *Humicola insolens* cell or a *Humicola lanuginosus* cell. In another preferred embodiment, the filamentous fungal host cell is a *Myceliophthora thermophila* cell. In a most preferred embodiment, the filamentous fungal host cell is a *Mucor miehei* cell. In a most preferred embodiment, the filamentous fungal host cell is a *Neurospora crassa* cell. In a most preferred embodiment, the filamentous fungal host cell is a *Penicillium purpurogenum* cell. In another most preferred embodiment, the filamentous fungal host cell is a *Thielavia terrestris* cell. In another most preferred embodiment, the *Trichoderma* cell is a *Trichoderma reesei* cell, a *Trichoderma viride* cell, a *Trichoderma longibrachiatum* cell, a *Trichoderma harzianum* cell, or a *Trichoderma koningii* cell. In a preferred embodiment, the fungal host cell is selected from an *A. nidulans* cell, an *A. niger* cell, an *A. oryzae* cell and an *A. sojae* cell. In a further preferred embodiment, the fungal host cell is an *A. nidulans* cell.

The recombinant fungal host cells of the present invention may further comprise one or more sequences which encode one or more factors that are advantageous in the expression of the protein or fragment thereof, for example, an activator (e.g., a trans-acting factor), a chaperone and a processing protease. The nucleic acids encoding one or more of these factors are preferably not operably linked to the nucleic acid encoding the protein or fragment thereof. An activator is a protein which activates transcription of a nucleic acid sequence encoding a polypeptide (Kudla *et al.*, *EMBO* 9:1355-1364(1990); Jarai and Buxton, *Current Genetics* 26:2238-244(1994); Verdier, *Yeast* 6:271-297(1990), all of which are herein incorporated by reference in their entirety). The nucleic acid sequence encoding an activator may be obtained from the genes encoding *Saccharomyces cerevisiae* heme activator protein 1 (hap1), *Saccharomyces cerevisiae* galactose metabolizing protein 4 (gal4) and *Aspergillus nidulans* ammonia regulation protein (areA). For further examples, see Verdier, *Yeast* 6:271-297 (1990);



MacKenzie *et al.*, *Journal of Gen. Microbiol.* 139:2295-2307 (1993), both of which are herein incorporated by reference in their entirety). A chaperone is a protein which assists another protein in folding properly (Hartl *et al.*, *TIBS* 19:20-25 (1994); Bergeron *et al.*, *TIBS* 19:124-128 (1994); Demolder *et al.*, *J. Biotechnology* 32:179-189 (1994); Craig, *Science* 260:1902-1903(1993); Gething and Sambrook, *Nature* 355:33-45 (1992); Puig and Gilbert, *J Biol. Chem.* 269:7764-7771 (1994); Wang and Tsou, *FASEB Journal* 7:1515-11157 (1993); Robinson *et al.*, *Bio/Technology* 1:381-384 (1994), all of which are herein incorporated by reference in their entirety). The nucleic acid sequence encoding a chaperone may be obtained from the genes encoding *Aspergillus oryzae* protein disulphide isomerase, *Saccharomyces cerevisiae* calnexin, *Saccharomyces cerevisiae* BiP/GRP78 and *Saccharomyces cerevisiae* Hsp70. For further examples, see Gething and Sambrook, *Nature* 355:33-45 (1992); Hartl *et al.*, *TIBS* 19:20-25 (1994). A processing protease is a protease that cleaves a propeptide to generate a mature biochemically active polypeptide (Enderlin and Ogrydziak, *Yeast* 10:67-79 (1994); Fuller *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 86:1434-1438 (1989); Julius *et al.*, *Cell* 37:1075-1089 (1984); Julius *et al.*, *Cell* 32:839-852 (1983), all of which are incorporated by reference in their entirety). The nucleic acid sequence encoding a processing protease may be obtained from the genes encoding *Aspergillus niger* Kex2, *Saccharomyces cerevisiae* dipeptidylaminopeptidase, *Saccharomyces cerevisiae* Kex2 and *Yarrowia lipolytica* dibasic processing endoprotease (xpr6). Any factor that is functional in the fungal host cell of choice may be used in the present invention.

Fungal cells may be transformed by a process involving protoplast formation, transformation of the protoplasts and regeneration of the cell wall in a manner known per se. Suitable procedures for transformation of *Aspergillus* host cells are described in EP 238 023 and

Yelton *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 81:1470-1474 (1984), both of which are herein incorporated by reference in their entirety. A suitable method of transforming *Fusarium* species is described by Malardier *et al.*, *Gene* 78:147-156 (1989), the entirety of which is herein incorporated by reference. Yeast may be transformed using the procedures described by Becker and Guarente, In: Abelson and Simon, (eds.), *Guide to Yeast Genetics and Molecular Biology, Methods Enzymol.* Volume 194, pp 182-187, Academic Press, Inc., New York; Ito *et al.*, *J. Bacteriology* 153:163 (1983); Hinnen *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 75:1920 (1978), all of which are herein incorporated by reference in their entirety.

The present invention also relates to methods of producing the protein or fragment thereof comprising culturing the recombinant fungal host cells under conditions conducive for expression of the protein or fragment thereof. The fungal cells of the present invention are cultivated in a nutrient medium suitable for production of the protein or fragment thereof using methods known in the art. For example, the cell may be cultivated by shake flask cultivation, small-scale or large-scale fermentation (including continuous, batch, fed-batch, or solid state fermentations) in laboratory or industrial fermentors performed in a suitable medium and under conditions allowing the protein or fragment thereof to be expressed and/or isolated. The cultivation takes place in a suitable nutrient medium comprising carbon and nitrogen sources and inorganic salts, using procedures known in the art (*see, e.g.*, Bennett and LaSure (eds.), *More Gene Manipulations in Fungi*, Academic Press, CA, (1991), the entirety of which is herein incorporated by reference). Suitable media are available from commercial suppliers or may be prepared according to published compositions (*e.g.*, in catalogues of the American Type Culture Collection, Manassas, VA). If the protein or fragment thereof is secreted into the nutrient

medium, a protein or fragment thereof can be recovered directly from the medium. If the protein or fragment thereof is not secreted, it is recovered from cell lysates.

The expressed protein or fragment thereof may be detected using methods known in the art that are specific for the particular protein or fragment. These detection methods may include the use of specific antibodies, formation of an enzyme product, or disappearance of an enzyme substrate. For example, if the protein or fragment thereof has enzymatic activity, an enzyme assay may be used. Alternatively, if polyclonal or monoclonal antibodies specific to the protein or fragment thereof are available, immunoassays may be employed using the antibodies to the protein or fragment thereof. The techniques of enzyme assay and immunoassay are well known to those skilled in the art.

The resulting protein or fragment thereof may be recovered by methods known in the arts. For example, the protein or fragment thereof may be recovered from the nutrient medium by conventional procedures including, but not limited to, centrifugation, filtration, extraction, spray-drying, evaporation, or precipitation. The recovered protein or fragment thereof may then be further purified by a variety of chromatographic procedures, e.g., ion exchange chromatography, gel filtration chromatography, affinity chromatography, or the like.

### **(c) Mammalian Constructs and Transformed Mammalian Cells**

The present invention also relates to methods for obtaining a recombinant mammalian host cell, comprising introducing into a mammalian host cell exogenous genetic material. The present invention also relates to a mammalian cell comprising a mammalian recombinant vector. The present invention also relates to methods for obtaining a recombinant mammalian host cell, comprising introducing into a mammalian cell exogenous genetic material.

Mammalian cell lines available as hosts for expression are known in the art and include many immortalized cell lines available from the American Type Culture Collection (ATCC, Manassas, VA), such as HeLa cells, Chinese hamster ovary (CHO) cells, baby hamster kidney (BHK) cells and a number of other cell lines. Suitable promoters for mammalian cells are also known in the art and include viral promoters such as that from Simian Virus 40 (SV40) (Fiers *et al.*, *Nature* 273:113 (1978), the entirety of which is herein incorporated by reference), Rous sarcoma virus (RSV), adenovirus (ADV) and bovine papilloma virus (BPV). Mammalian cells may also require terminator sequences and poly-A addition sequences. Enhancer sequences which increase expression may also be included and sequences which promote amplification of the gene may also be desirable (for example methotrexate resistance genes).

Vectors suitable for replication in mammalian cells may include viral replicons, or sequences which insure integration of the appropriate sequences encoding HCV epitopes into the host genome. For example, another vector used to express foreign DNA is vaccinia virus. In this case, for example, a nucleic acid molecule encoding a protein or fragment thereof is inserted into the vaccinia genome. Techniques for the insertion of foreign DNA into the vaccinia virus genome are known in the art and may utilize, for example, homologous recombination. Such heterologous DNA is generally inserted into a gene which is non-essential to the virus, for example, the thymidine kinase gene (tk), which also provides a selectable marker. Plasmid vectors that greatly facilitate the construction of recombinant viruses have been described (*see*, for example, Mackett *et al.*, *J Virol.* 49:857 (1984); Chakrabarti *et al.*, *Mol. Cell. Biol.* 5:3403 (1985); Moss, In: *Gene Transfer Vectors For Mammalian Cells* (Miller and Calos, eds., Cold Spring Harbor Laboratory, N.Y., p. 10, (1987); all of which are herein incorporated by reference

in their entirety). Expression of the HCV polypeptide then occurs in cells or animals which are infected with the live recombinant vaccinia virus.

The sequence to be integrated into the mammalian sequence may be introduced into the primary host by any convenient means, which includes calcium precipitated DNA, spheroplast fusion, transformation, electroporation, biolistics, lipofection, microinjection, or other convenient means. Where an amplifiable gene is being employed, the amplifiable gene may serve as the selection marker for selecting hosts into which the amplifiable gene has been introduced. Alternatively, one may include with the amplifiable gene another marker, such as a drug resistance marker, e.g. neomycin resistance (G418 in mammalian cells), hygromycin in resistance etc., or an auxotrophy marker (HIS3, TRP1, LEU2, URA3, ADE2, LYS2, etc.) for use in yeast cells.

Depending upon the nature of the modification and associated targeting construct, various techniques may be employed for identifying targeted integration. Conveniently, the DNA may be digested with one or more restriction enzymes and the fragments probed with an appropriate DNA fragment which will identify the properly sized restriction fragment associated with integration.

One may use different promoter sequences, enhancer sequences, or other sequence which will allow for enhanced levels of expression in the expression host. Thus, one may combine an enhancer from one source, a promoter region from another source, a 5'- noncoding region upstream from the initiation methionine from the same or different source as the other sequences and the like. One may provide for an intron in the non-coding region with appropriate splice sites or for an alternative 3'- untranslated sequence or polyadenylation site. Depending upon the particular purpose of the modification, any of these sequences may be introduced, as desired.

Where selection is intended, the sequence to be integrated will have with it a marker gene, which allows for selection. The marker gene may conveniently be downstream from the target gene and may include resistance to a cytotoxic agent, e.g. antibiotics, heavy metals, or the like, resistance or susceptibility to HAT, gancyclovir, etc., complementation to an auxotrophic host, particularly by using an auxotrophic yeast as the host for the subject manipulations, or the like. The marker gene may also be on a separate DNA molecule, particularly with primary mammalian cells. Alternatively, one may screen the various transformants, due to the high efficiency of recombination in yeast, by using hybridization analysis, PCR, sequencing, or the like.

For homologous recombination, constructs can be prepared where the amplifiable gene will be flanked, normally on both sides with DNA homologous with the DNA of the target region. Depending upon the nature of the integrating DNA and the purpose of the integration, the homologous DNA will generally be within 100kb, usually 50kb, preferably about 25kb, of the transcribed region of the target gene, more preferably within 2kb of the target gene. Where modeling of the gene is intended, homology will usually be present proximal to the site of the mutation. The homologous DNA may include the 5'-upstream region outside of the transcriptional regulatory region or comprising any enhancer sequences, transcriptional initiation sequences, adjacent sequences, or the like. The homologous region may include a portion of the coding region, where the coding region may be comprised only of an open reading frame or combination of exons and introns. The homologous region may comprise all or a portion of an intron, where all or a portion of one or more exons may also be present. Alternatively, the homologous region may comprise the 3'-region, so as to comprise all or a portion of the transcriptional termination region, or the region 3' of this region. The homologous regions may

extend over all or a portion of the target gene or be outside the target gene comprising all or a portion of the transcriptional regulatory regions and/or the structural gene.

The integrating constructs may be prepared in accordance with conventional ways, where sequences may be synthesized, isolated from natural sources, manipulated, cloned, ligated, subjected to in vitro mutagenesis, primer repair, or the like. At various stages, the joined sequences may be cloned and analyzed by restriction analysis, sequencing, or the like. Usually during the preparation of a construct where various fragments are joined, the fragments, intermediate constructs and constructs will be carried on a cloning vector comprising a replication system functional in a prokaryotic host, e.g., *E. coli* and a marker for selection, e.g., biocide resistance, complementation to an auxotrophic host, etc. Other functional sequences may also be present, such as polylinkers, for ease of introduction and excision of the construct or portions thereof, or the like. A large number of cloning vectors are available such as pBR322, the pUC series, etc. These constructs may then be used for integration into the primary mammalian host.

In the case of the primary mammalian host, a replicating vector may be used. Usually, such vector will have a viral replication system, such as SV40, bovine papilloma virus, adenovirus, or the like. The linear DNA sequence vector may also have a selectable marker for identifying transfected cells. Selectable markers include the neo gene, allowing for selection with G418, the herpes tk gene for selection with HAT medium, the gpt gene with mycophenolic acid, complementation of an auxotrophic host, etc.

The vector may or may not be capable of stable maintenance in the host. Where the vector is capable of stable maintenance, the cells will be screened for homologous integration of the vector into the genome of the host, where various techniques for curing the cells may be

employed. Where the vector is not capable of stable maintenance, for example, where a temperature sensitive replication system is employed, one may change the temperature from the permissive temperature to the non-permissive temperature, so that the cells may be cured of the vector. In this case, only those cells having integration of the construct comprising the amplifiable gene and, when present, the selectable marker, will be able to survive selection.

Where a selectable marker is present, one may select for the presence of the targeting construct by means of the selectable marker. Where the selectable marker is not present, one may select for the presence of the construct by the amplifiable gene. For the neo gene or the herpes tk gene, one could employ a medium for growth of the transformants of about 0.1-1 mg/ml of G418 or may use HAT medium, respectively. Where DHFR is the amplifiable gene, the selective medium may include from about 0.01-0.5  $\mu$ M of methotrexate or be deficient in glycine-hypoxanthine-thymidine and have dialysed serum (GHT media).

The DNA can be introduced into the expression host by a variety of techniques that include calcium phosphate/DNA co-precipitates, microinjection of DNA into the nucleus, electroporation, yeast protoplast fusion with intact cells, transfection, polycations, e.g., polybrene, polyornithine, etc., or the like. The DNA may be single or double stranded DNA, linear or circular. The various techniques for transforming mammalian cells are well known (see Keown *et al.*, *Methods Enzymol.* (1989); Keown *et al.*, *Methods Enzymol.* 185:527-537 (1990); Mansour *et al.*, *Nature* 336:348-352, (1988); all of which are herein incorporated by reference in their entirety).

#### **(d) Insect Constructs and Transformed Insect Cells**

The present invention also relates to an insect recombinant vectors comprising exogenous genetic material. The present invention also relates to an insect cell comprising an insect



recombinant vector. The present invention also relates to methods for obtaining a recombinant insect host cell, comprising introducing into an insect cell exogenous genetic material.

The insect recombinant vector may be any vector which can be conveniently subjected to recombinant DNA procedures and can bring about the expression of the nucleic acid sequence. The choice of a vector will typically depend on the compatibility of the vector with the insect host cell into which the vector is to be introduced. The vector may be a linear or a closed circular plasmid. The vector system may be a single vector or plasmid or two or more vectors or plasmids which together contain the total DNA to be introduced into the genome of the insect host. In addition, the insect vector may be an expression vector. Nucleic acid molecules can be suitably inserted into a replication vector for expression in the insect cell under a suitable promoter for insect cells. Many vectors are available for this purpose and selection of the appropriate vector will depend mainly on the size of the nucleic acid molecule to be inserted into the vector and the particular host cell to be transformed with the vector. Each vector contains various components depending on its function (amplification of DNA or expression of DNA) and the particular host cell with which it is compatible. The vector components for insect cell transformation generally include, but are not limited to, one or more of the following: a signal sequence, origin of replication, one or more marker genes and an inducible promoter.

The insect vector may be an autonomously replicating vector, *i.e.*, a vector which exists as an extrachromosomal entity, the replication of which is independent of chromosomal replication, *e.g.*, a plasmid, an extrachromosomal element, a minichromosome, or an artificial chromosome. The vector may contain any means for assuring self-replication. Alternatively, the vector may be one which, when introduced into the insect cell, is integrated into the genome and replicated together with the chromosome(s) into which it has been integrated. For integration,

the vector may rely on the nucleic acid sequence of the vector for stable integration of the vector into the genome by homologous or nonhomologous recombination. Alternatively, the vector may contain additional nucleic acid sequences for directing integration by homologous recombination into the genome of the insect host. The additional nucleic acid sequences enable the vector to be integrated into the host cell genome at a precise location(s) in the chromosome(s). To increase the likelihood of integration at a precise location, there should be preferably two nucleic acid sequences which individually contain a sufficient number of nucleic acids, preferably 400bp to 1500bp, more preferably 800bp to 1000bp, which are highly homologous with the corresponding target sequence to enhance the probability of homologous recombination. These nucleic acid sequences may be any sequence that is homologous with a target sequence in the genome of the insect host cell and, furthermore, may be non-encoding or encoding sequences.

Baculovirus expression vectors (BEVs) have become important tools for the expression of foreign genes, both for basic research and for the production of proteins with direct clinical applications in human and veterinary medicine (Doerfler, *Curr. Top. Microbiol. Immunol.* 131:51-68 (1968); Luckow and Summers, *Bio/Technology* 6:47-55 (1988a); Miller, *Annual Review of Microbiol.* 42:177-199 (1988); Summers, *Curr. Comm. Molecular Biology*, Cold Spring Harbor Press, Cold Spring Harbor, N.Y. (1988); all of which are herein incorporated by reference in their entirety). BEVs are recombinant insect viruses in which the coding sequence for a chosen foreign gene has been inserted behind a baculovirus promoter in place of the viral gene, e.g., polyhedrin (Smith and Summers, U.S. Pat. No., 4,745,051, the entirety of which is incorporated herein by reference).

The use of baculovirus vectors relies upon the host cells being derived from *Lepidopteran* insects such as *Spodoptera frugiperda* or *Trichoplusia ni*. The preferred *Spodoptera frugiperda* cell line is the cell line Sf9. The *Spodoptera frugiperda* Sf9 cell line was obtained from American Type Culture Collection (Manassas, VA.) and is assigned accession number ATCC CRL 1711 (Summers and Smith, *A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures*, Texas Ag. Exper. Station Bulletin No. 1555 (1988), the entirety of which is herein incorporated by reference). Other insect cell systems, such as the silkworm *B. mori* may also be used.

The proteins expressed by the BEVs are, therefore, synthesized, modified and transported in host cells derived from *Lepidopteran* insects. Most of the genes that have been inserted and produced in the baculovirus expression vector system have been derived from vertebrate species. Other baculovirus genes in addition to the polyhedrin promoter may be employed to advantage in a baculovirus expression system. These include immediate-early (alpha), delayed-early ( $\beta$ ), late ( $\gamma$ ), or very late (delta), according to the phase of the viral infection during which they are expressed. The expression of these genes occurs sequentially, probably as the result of a "cascade" mechanism of transcriptional regulation. (Guarino and Summers, *J. Virol.* 57:563-571 (1986); Guarino and Summers, *J. Virol.* 61:2091-2099 (1987); Guarino and Summers, *Virol.* 162:444-451 (1988); all of which are herein incorporated by reference in their entirety).

Insect recombinant vectors are useful as intermediates for the infection or transformation of insect cell systems. For example, an insect recombinant vector containing a nucleic acid molecule encoding a baculovirus transcriptional promoter followed downstream by an insect signal DNA sequence is capable of directing the secretion of the desired biologically active protein from the insect cell. The vector may utilize a baculovirus transcriptional promoter region

derived from any of the over 500 baculoviruses generally infecting insects, such as for example the Orders *Lepidoptera*, *Diptera*, *Orthoptera*, *Coleoptera* and *Hymenoptera*, including for example but not limited to the viral DNAs of *Autographa californica* MNPV, *Bombyx mori* NPV, *Trichoplusia ni* MNPV, *Rachiplusia ou* MNPV or *Galleria mellonella* MNPV, wherein said baculovirus transcriptional promoter is a baculovirus immediate-early gene IEl or IEN promoter; an immediate-early gene in combination with a baculovirus delayed-early gene promoter region selected from the group consisting of 39K and a *HindIII-k* fragment delayed-early gene; or a baculovirus late gene promoter. The immediate-early or delayed-early promoters can be enhanced with transcriptional enhancer elements. The insect signal DNA sequence may code for a signal peptide of a *Lepidopteran* adipokinetic hormone precursor or a signal peptide of the *Manduca sexta* adipokinetic hormone precursor (Summers, U.S. Patent No. 5,155,037; the entirety of which is herein incorporated by reference). Other insect signal DNA sequences include a signal peptide of the *Orthoptera Schistocerca gregaria* locust adipokinetic hormone precursor and the *Drosophila melanogaster* cuticle genes CP1, CP2, CP3 or CP4 or for an insect signal peptide having substantially a similar chemical composition and function (Summers, U.S. Patent No. 5,155,037).

Insect cells are distinctly different from animal cells. Insects have a unique life cycle and have distinct cellular properties such as the lack of intracellular plasminogen activators in which are present in vertebrate cells. Another difference is the high expression levels of protein products ranging from 1 to greater than 500 mg/liter and the ease at which cDNA can be cloned into cells (Frasier, *In Vitro Cell. Dev. Biol.* 25:225 (1989); Summers and Smith, In: *A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures*, Texas Ag. Exper. Station Bulletin No. 1555 (1988), both of which are incorporated by reference in their entirety).

Recombinant protein expression in insect cells is achieved by viral infection or stable transformation. For viral infection, the desired gene is cloned into baculovirus at the site of the wild-type polyhedron gene (Webb and Summers, *Technique* 2:173 (1990); Bishop and Posse, *Adv. Gene Technol.* 1:55 (1990); both of which are incorporated by reference in their entirety). The polyhedron gene is a component of a protein coat in occlusions which encapsulate virus particles. Deletion or insertion in the polyhedron gene results the failure to form occlusion bodies. Occlusion negative viruses are morphologically different from occlusion positive viruses and enable one skilled in the art to identify and purify recombinant viruses.

The vectors of present invention preferably contain one or more selectable markers which permit easy selection of transformed cells. A selectable marker is a gene the product of which provides, for example biocide or viral resistance, resistance to heavy metals, prototrophy to auxotrophs and the like. Selection may be accomplished by co-transformation, *e.g.*, as described in WO 91/17243, a nucleic acid sequence of the present invention may be operably linked to a suitable promoter sequence. The promoter sequence is a nucleic acid sequence which is recognized by the insect host cell for expression of the nucleic acid sequence. The promoter sequence contains transcription and translation control sequences which mediate the expression of the protein or fragment thereof. The promoter may be any nucleic acid sequence which shows transcriptional activity in the insect host cell of choice and may be obtained from genes encoding polypeptides either homologous or heterologous to the host cell.

For example, a nucleic acid molecule encoding a protein or fragment thereof may also be operably linked to a suitable leader sequence. A leader sequence is a nontranslated region of a mRNA which is important for translation by the fungal host. The leader sequence is operably linked to the 5' terminus of the nucleic acid sequence encoding the protein or fragment thereof.

The leader sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any leader sequence which is functional in the insect host cell of choice may be used in the present invention.

A polyadenylation sequence may also be operably linked to the 3' terminus of the nucleic acid sequence of the present invention. The polyadenylation sequence is a sequence which when transcribed is recognized by the insect host to add polyadenosine residues to transcribed mRNA. The polyadenylation sequence may be native to the nucleic acid sequence encoding the protein or fragment thereof or may be obtained from foreign sources. Any polyadenylation sequence which is functional in the fungal host of choice may be used in the present invention.

To avoid the necessity of disrupting the cell to obtain the protein or fragment thereof and to minimize the amount of possible degradation of the expressed polypeptide within the cell, it is preferred that expression of the polypeptide gene gives rise to a product secreted outside the cell.

To this end, the protein or fragment thereof of the present invention may be linked to a signal peptide linked to the amino terminus of the protein or fragment thereof. A signal peptide is an amino acid sequence which permits the secretion of the protein or fragment thereof from the insect host into the culture medium. The signal peptide may be native to the protein or fragment thereof of the invention or may be obtained from foreign sources. The 5' end of the coding sequence of the nucleic acid sequence of the present invention may inherently contain a signal peptide coding region naturally linked in translation reading frame with the segment of the coding region which encodes the secreted protein or fragment thereof.

At present, a mode of achieving secretion of a foreign gene product in insect cells is by way of the foreign gene's native signal peptide. Because the foreign genes are usually from non-insect organisms, their signal sequences may be poorly recognized by insect cells and hence,

levels of expression may be suboptimal. However, the efficiency of expression of foreign gene products seems to depend primarily on the characteristics of the foreign protein. On average, nuclear localized or non-structural proteins are most highly expressed, secreted proteins are intermediate and integral membrane proteins are the least expressed. One factor generally affecting the efficiency of the production of foreign gene products in a heterologous host system is the presence of native signal sequences (also termed presequences, targeting signals, or leader sequences) associated with the foreign gene. The signal sequence is generally coded by a DNA sequence immediately following (5' to 3') the translation start site of the desired foreign gene.

The expression dependence on the type of signal sequence associated with a gene product can be represented by the following example: If a foreign gene is inserted at a site downstream from the translational start site of the baculovirus polyhedrin gene so as to produce a fusion protein (containing the N-terminus of the polyhedrin structural gene), the fused gene is highly expressed. But less expression is achieved when a foreign gene is inserted in a baculovirus expression vector immediately following the transcriptional start site and totally replacing the polyhedrin structural gene.

Insertions into the region -50 to -1 significantly alter (reduce) steady state transcription which, in turn, reduces translation of the foreign gene product. Use of the pVL941 vector optimizes transcription of foreign genes to the level of the polyhedrin gene transcription. Even though the transcription of a foreign gene may be optimal, optimal translation may vary because of several factors involving processing: signal peptide recognition, mRNA and ribosome binding, glycosylation, disulfide bond formation, sugar processing, oligomerization, for example.

The properties of the insect signal peptide are expected to be more optimal for the efficiency of the translation process in insect cells than those from vertebrate proteins. This

phenomenon can generally be explained by the fact that proteins secreted from cells are synthesized as precursor molecules containing hydrophobic N-terminal signal peptides. The signal peptides direct transport of the select protein to its target membrane and are then cleaved by a peptidase on the membrane, such as the endoplasmic reticulum, when the protein passes through it.

Another exemplary insect signal sequence is the sequence encoding for *Drosophila* cuticle proteins such as CP1, CP2, CP3 or CP4 (Summers, U.S. Patent No. 5,278,050; the entirety of which is herein incorporated by reference). Most of a 9kb region of the *Drosophila* genome containing genes for the cuticle proteins has been sequenced. Four of the five cuticle genes contains a signal peptide coding sequence interrupted by a short intervening sequence (about 60 base pairs) at a conserved site. Conserved sequences occur in the 5' mRNA untranslated region, in the adjacent 35 base pairs of upstream flanking sequence and at -200 base pairs from the mRNA start position in each of the cuticle genes.

Standard methods of insect cell culture, cotransfection and preparation of plasmids are set forth in Summers and Smith (Summers and Smith, *A Manual of Methods for Baculovirus Vectors and Insect Cell Culture Procedures*, Texas Agricultural Experiment Station Bulletin No. 1555, Texas A&M University (1987)). Procedures for the cultivation of viruses and cells are described in Volkman and Summers, *J. Virol* 19:820-832 (1975) and Volkman *et al.*, *J. Virol* 19:820-832 (1976); both of which are herein incorporated by reference in their entirety.

#### **(e) Bacterial Constructs and Transformed Bacterial Cells**

The present invention also relates to a bacterial recombinant vector comprising exogenous genetic material. The present invention also relates to a bacteria cell comprising a bacterial recombinant vector. The present invention also relates to methods for obtaining a



recombinant bacteria host cell, comprising introducing into a bacterial host cell exogenous genetic material.

The bacterial recombinant vector may be any vector which can be conveniently subjected to recombinant DNA procedures. The choice of a vector will typically depend on the compatibility of the vector with the bacterial host cell into which the vector is to be introduced. The vector may be a linear or a closed circular plasmid. The vector system may be a single vector or plasmid or two or more vectors or plasmids which together contain the total DNA to be introduced into the genome of the bacterial host. In addition, the bacterial vector may be an expression vector. Nucleic acid molecules encoding protein homologues or fragments thereof can, for example, be suitably inserted into a replicable vector for expression in the bacterium under the control of a suitable promoter for bacteria. Many vectors are available for this purpose and selection of the appropriate vector will depend mainly on the size of the nucleic acid to be inserted into the vector and the particular host cell to be transformed with the vector. Each vector contains various components depending on its function (amplification of DNA or expression of DNA) and the particular host cell with which it is compatible. The vector components for bacterial transformation generally include, but are not limited to, one or more of the following: a signal sequence, an origin of replication, one or more marker genes and an inducible promoter.

In general, plasmid vectors containing replicon and control sequences that are derived from species compatible with the host cell are used in connection with bacterial hosts. The vector ordinarily carries a replication site, as well as marking sequences that are capable of providing phenotypic selection in transformed cells. For example, *E. coli* is typically transformed using pBR322, a plasmid derived from an *E. coli* species (see, e.g., Bolivar *et al.*,

*Gene* 2:95 (1977); the entirety of which is herein incorporated by reference). pBR322 contains genes for ampicillin and tetracycline resistance and thus provides easy means for identifying transformed cells. The pBR322 plasmid, or other microbial plasmid or phage, also generally contains, or is modified to contain, promoters that can be used by the microbial organism for expression of the selectable marker genes.

Nucleic acid molecules encoding protein or fragments thereof may be expressed not only directly, but also as a fusion with another polypeptide, preferably a signal sequence or other polypeptide having a specific cleavage site at the N-terminus of the mature polypeptide. In general, the signal sequence may be a component of the vector, or it may be a part of the polypeptide DNA that is inserted into the vector. The heterologous signal sequence selected should be one that is recognized and processed (i.e., cleaved by a signal peptidase) by the host cell. For bacterial host cells that do not recognize and process the native polypeptide signal sequence, the signal sequence is substituted by a bacterial signal sequence selected, for example, from the group consisting of the alkaline phosphatase, penicillinase, lpp, or heat-stable enterotoxin II leaders.

Both expression and cloning vectors contain a nucleic acid sequence that enables the vector to replicate in one or more selected host cells. Generally, in cloning vectors this sequence is one that enables the vector to replicate independently of the host chromosomal DNA and includes origins of replication or autonomously replicating sequences. Such sequences are well known for a variety of bacteria. The origin of replication from the plasmid pBR322 is suitable for most Gram-negative bacteria.

Expression and cloning vectors also generally contain a selection gene, also termed a selectable marker. This gene encodes a protein necessary for the survival or growth of

transformed host cells grown in a selective culture medium. Host cells not transformed with the vector containing the selection gene will not survive in the culture medium. Typical selection genes encode proteins that (a) confer resistance to antibiotics or other toxins, e.g., ampicillin, neomycin, methotrexate, or tetracycline, (b) complement auxotrophic deficiencies, or (c) supply critical nutrients not available from complex media, e.g., the gene encoding D-alanine racemase for *Bacilli*. One example of a selection scheme utilizes a drug to arrest growth of a host cell. Those cells that are successfully transformed with a heterologous protein homologue or fragment thereof produce a protein conferring drug resistance and thus survive the selection regimen.

The expression vector for producing a protein or fragment thereof can also contains an inducible promoter that is recognized by the host bacterial organism and is operably linked to the nucleic acid encoding, for example, the nucleic acid molecule encoding the protein homologue or fragment thereof of interest. Inducible promoters suitable for use with bacterial hosts include the  $\beta$ -lactamase and lactose promoter systems (Chang *et al.*, *Nature* 275:615 (1978); Goeddel *et al.*, *Nature* 281:544 (1979); both of which are herein incorporated by reference in their entirety), the arabinose promoter system (Guzman *et al.*, *J. Bacteriol.* 174:7716-7728 (1992); the entirety of which is herein incorporated by reference), alkaline phosphatase, a tryptophan (trp) promoter system (Goeddel, *Nucleic Acids Res.* 8:4057 (1980); EP 36,776; both of which are herein incorporated by reference in their entirety) and hybrid promoters such as the tac promoter (deBoer *et al.*, *Proc. Natl. Acad. Sci. (USA)* 80:21-25 (1983); the entirety of which is herein incorporated by reference). However, other known bacterial inducible promoters are suitable (Siebenlist *et al.*, *Cell* 20:269 (1980); the entirety of which is herein incorporated by reference).

Promoters for use in bacterial systems also generally contain a Shine-Dalgarno (S.D.) sequence operably linked to the DNA encoding the polypeptide of interest. The promoter can be

removed from the bacterial source DNA by restriction enzyme digestion and inserted into the vector containing the desired DNA.

Construction of suitable vectors containing one or more of the above-listed components employs standard ligation techniques. Isolated plasmids or DNA fragments are cleaved, tailored and re-ligated in the form desired to generate the plasmids required. Examples of available bacterial expression vectors include, but are not limited to, the multifunctional *E. coli* cloning and expression vectors such as Bluescript™ (Stratagene, La Jolla, CA), in which, for example, encoding an *A. nidulans* protein homologue or fragment thereof homologue, may be ligated into the vector in frame with sequences for the amino-terminal Met and the subsequent 7 residues of β-galactosidase so that a hybrid protein is produced; pIN vectors (Van Heeke and Schuster, *J. Biol. Chem.* 264:5503-5509 (1989), the entirety of which is herein incorporated by reference); and the like. pGEX vectors (Promega, Madison Wisconsin U.S.A.) may also be used to express foreign polypeptides as fusion proteins with glutathione S-transferase (GST). In general, such fusion proteins are soluble and can easily be purified from lysed cells by adsorption to glutathione-agarose beads followed by elution in the presence of free glutathione. Proteins made in such systems are designed to include heparin, thrombin or factor XA protease cleavage sites so that the cloned polypeptide of interest can be released from the GST moiety at will.

Suitable host bacteria for a bacterial vector include archaebacteria and eubacteria, especially eubacteria and most preferably *Enterobacteriaceae*. Examples of useful bacteria include *Escherichia*, *Enterobacter*, *Azotobacter*, *Erwinia*, *Bacillus*, *Pseudomonas*, *Klebsiella*, *Proteus*, *Salmonella*, *Serratia*, *Shigella*, *Rhizobia*, *Vitreoscilla* and *Paracoccus*. Suitable *E. coli* hosts include *E. coli* W3110 (American Type Culture Collection (ATCC) 27,325, Manassas,

Virginia U.S.A.), *E. coli* 294 (ATCC 31,446), *E. coli* B and *E. coli* X1776 (ATCC 31,537).

These examples are illustrative rather than limiting. Mutant cells of any of the above-mentioned bacteria may also be employed. It is, of course, necessary to select the appropriate bacteria taking into consideration replicability of the replicon in the cells of a bacterium. For example, *E. coli*, *Serratia*, or *Salmonella* species can be suitably used as the host when well known plasmids such as pBR322, pBR325, pACYC177, or pKN410 are used to supply the replicon. *E. coli* strain W3110 is a preferred host or parent host because it is a common host strain for recombinant DNA product fermentations. Preferably, the host cell should secrete minimal amounts of proteolytic enzymes.

Host cells are transfected and preferably transformed with the above-described vectors and cultured in conventional nutrient media modified as appropriate for inducing promoters, selecting transformants, or amplifying the genes encoding the desired sequences.

Numerous methods of transfection are known to the ordinarily skilled artisan, for example, calcium phosphate and electroporation. Depending on the host cell used, transformation is done using standard techniques appropriate to such cells. The calcium treatment employing calcium chloride, as described in section 1.82 of Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, New York: Cold Spring Harbor Laboratory Press, (1989), is generally used for bacterial cells that contain substantial cell-wall barriers. Another method for transformation employs polyethylene glycol/DMSO, as described in Chung and Miller (Chung and Miller, *Nucleic Acids Res.* 16:3580 (1988); the entirety of which is herein incorporated by reference). Yet another method is the use of the technique termed electroporation.

Bacterial cells used to produce the polypeptide of interest for purposes of this invention are cultured in suitable media in which the promoters for the nucleic acid encoding the heterologous polypeptide can be artificially induced as described generally, e.g., in Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, New York: Cold Spring Harbor Laboratory Press, (1989). Examples of suitable media are given in U.S. Pat. Nos. 5,304,472 and 5,342,763; both of which are incorporated by reference in their entirety.

In addition to the above discussed procedures, practitioners are familiar with the standard resource materials which describe specific conditions and procedures for the construction, manipulation and isolation of macromolecules (e.g., DNA molecules, plasmids, etc.), generation of recombinant organisms and the screening and isolating of clones, (see for example, Sambrook *et al.*, *Molecular Cloning: A Laboratory Manual*, Cold Spring Harbor Press (1989); Mailga *et al.*, *Methods in Plant Molecular Biology*, Cold Spring Harbor Press (1995), the entirety of which is herein incorporated by reference; Birren *et al.*, *Genome Analysis: Analyzing DNA*, 1, Cold Spring Harbor, New York, the entirety of which is herein incorporated by reference).

#### **(f) Computer Readable Media**

The nucleotide sequence provided in SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof, or complement thereof, or a nucleotide sequence at least 90% identical, preferably 95%, identical even more preferably 99% or 100% identical to the sequence provided in SEQ ID NO: 1 through SEQ ID NO: 3204 or fragment thereof, or complement thereof, can be “provided” in a variety of mediums to facilitate use. Such a medium can also provide a subset thereof in a form that allows a skilled artisan to examine the sequences.

A preferred subset of nucleotide sequences are those nucleic acid sequences that encode a maize or soybean methionine adenosyltransferase enzyme or complement thereof or fragment of

either, a nucleic acid molecule that encodes a maize or soybean S-adenosylmethionine  
 decarboxylase enzyme or complement thereof or fragment of either, a nucleic acid molecule that  
 encodes a maize or soybean aspartate kinase enzyme or complement thereof or fragment of  
 either, a nucleic acid molecule that encodes a maize or soybean aspartate-semialdehyde  
 dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that  
 encodes a maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof  
 or fragment of either, a nucleic acid molecule that encodes a maize or soybean cystathionine  $\beta$ -  
 lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes  
 a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase  
 enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a  
 maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of  
 either, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase  
 enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a  
 maize or a soybean cytsathionine  $\gamma$ -lyase enzyme or complement thereof or fragment of either  
 and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase  
 enzyme or complement thereof or fragment of either.

A further preferred subset of nucleic acid sequences is where the subset of sequences is  
 two proteins or fragments thereof, more preferably three proteins or fragments thereof and even  
 more preferable four proteins or fragments thereof, these nucleic acid sequences are selected  
 from the group that comprises a maize or soybean methionine adenosyltransferase enzyme or  
 complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or  
 soybean S-adenosylmethionine decarboxylase enzyme or complement thereof or fragment of  
 either, a nucleic acid molecule that encodes a maize or soybean aspartate kinase enzyme or

complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean aspartate-semialdehyde dehydrogenase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean *O*-succinylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean cystathionine  $\beta$ -lyase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or soybean 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean adenosylhomocysteinase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cystathionine  $\beta$ -synthase enzyme or complement thereof or fragment of either, a nucleic acid molecule that encodes a maize or a soybean cytsathionine  $\gamma$ -lyase enzyme or complement thereof or fragment of either and a nucleic acid molecule that encodes a maize or a soybean *O*-acetylhomoserine (thiol)-lyase enzyme or complement thereof or fragment of either.

In one application of this embodiment, a nucleotide sequence of the present invention can be recorded on computer readable media. As used herein, "computer readable media" refers to any medium that can be read and accessed directly by a computer. Such media include, but are not limited to: magnetic storage media, such as floppy discs, hard disc, storage medium and magnetic tape; optical storage media such as CD-ROM; electrical storage media such as RAM and ROM; and hybrids of these categories such as magnetic/optical storage media. A skilled artisan can readily appreciate how any of the presently known computer readable mediums can be used to create a manufacture comprising computer readable medium having recorded thereon a nucleotide sequence of the present invention.



As used herein, "recorded" refers to a process for storing information on computer readable medium. A skilled artisan can readily adopt any of the presently known methods for recording information on computer readable medium to generate media comprising the nucleotide sequence information of the present invention. A variety of data storage structures are available to a skilled artisan for creating a computer readable medium having recorded thereon a nucleotide sequence of the present invention. The choice of the data storage structure will generally be based on the means chosen to access the stored information. In addition, a variety of data processor programs and formats can be used to store the nucleotide sequence information of the present invention on computer readable medium. The sequence information can be represented in a word processing text file, formatted in commercially-available software such as WordPerfect and Microsoft Word, or represented in the form of an ASCII file, stored in a database application, such as DB2, Sybase, Oracle, or the like. A skilled artisan can readily adapt any number of data processor structuring formats (e.g. text file or database) in order to obtain computer readable medium having recorded thereon the nucleotide sequence information of the present invention.

By providing one or more of nucleotide sequences of the present invention, a skilled artisan can routinely access the sequence information for a variety of purposes. Computer software is publicly available which allows a skilled artisan to access sequence information provided in a computer readable medium. The examples which follow demonstrate how software which implements the BLAST (Altschul *et al.*, *J. Mol. Biol.* 215:403-410 (1990), the entirety of which is herein incorporated by reference) and BLAZE (Brutlag *et al.*, *Comp. Chem.* 17:203-207 (1993), the entirety of which is herein incorporated by reference) search algorithms on a Sybase system can be used to identify open reading frames (ORFs) within the genome that

contain homology to ORFs or proteins from other organisms. Such ORFs are protein-encoding fragments within the sequences of the present invention and are useful in producing commercially important proteins such as enzymes used in amino acid biosynthesis, metabolism, transcription, translation, RNA processing, nucleic acid and a protein degradation, protein modification and DNA replication, restriction, modification, recombination and repair.

The present invention further provides systems, particularly computer-based systems, which contain the sequence information described herein. Such systems are designed to identify commercially important fragments of the nucleic acid molecule of the present invention. As used herein, “a computer-based system” refers to the hardware means, software means and data storage means used to analyze the nucleotide sequence information of the present invention. The minimum hardware means of the computer-based systems of the present invention comprises a central processing unit (CPU), input means, output means and data storage means. A skilled artisan can readily appreciate that any one of the currently available computer-based system are suitable for use in the present invention.

As indicated above, the computer-based systems of the present invention comprise a data storage means having stored therein a nucleotide sequence of the present invention and the necessary hardware means and software means for supporting and implementing a search means. As used herein, “data storage means” refers to memory that can store nucleotide sequence information of the present invention, or a memory access means which can access manufactures having recorded thereon the nucleotide sequence information of the present invention. As used herein, “search means” refers to one or more programs which are implemented on the computer-based system to compare a target sequence or target structural motif with the sequence information stored within the data storage means. Search means are used to identify fragments

or regions of the sequence of the present invention that match a particular target sequence or target motif. A variety of known algorithms are disclosed publicly and a variety of commercially available software for conducting search means are available can be used in the computer-based systems of the present invention. Examples of such software include, but are not limited to, MacPattern (EMBL), BLASTIN and BLASTIX (NCBIA). One of the available algorithms or implementing software packages for conducting homology searches can be adapted for use in the present computer-based systems.

The most preferred sequence length of a target sequence is from about 10 to 100 amino acids or from about 30 to 300 nucleotide residues. However, it is well recognized that during searches for commercially important fragments of the nucleic acid molecules of the present invention, such as sequence fragments involved in gene expression and protein processing, may be of shorter length.

As used herein, "a target structural motif," or "target motif," refers to any rationally selected sequence or combination of sequences in which the sequences the sequence(s) are chosen based on a three-dimensional configuration which is formed upon the folding of the target motif. There are a variety of target motifs known in the art. Protein target motifs include, but are not limited to, enzymatic active sites and signal sequences. Nucleic acid target motifs include, but are not limited to, promoter sequences, *cis* elements, hairpin structures and inducible expression elements (protein binding sequences).

Thus, the present invention further provides an input means for receiving a target sequence, a data storage means for storing the target sequences of the present invention sequence identified using a search means as described above and an output means for outputting the identified homologous sequences. A variety of structural formats for the input and output means

can be used to input and output information in the computer-based systems of the present invention. A preferred format for an output means ranks fragments of the sequence of the present invention by varying degrees of homology to the target sequence or target motif. Such presentation provides a skilled artisan with a ranking of sequences which contain various amounts of the target sequence or target motif and identifies the degree of homology contained in the identified fragment.

A variety of comparing means can be used to compare a target sequence or target motif with the data storage means to identify sequence fragments sequence of the present invention. For example, implementing software which implement the BLAST and BLAZE algorithms (Altschul *et al.*, *J. Mol. Biol.* 215:403-410 (1990)) can be used to identify open frames within the nucleic acid molecules of the present invention. A skilled artisan can readily recognize that any one of the publicly available homology search programs can be used as the search means for the computer-based systems of the present invention.

Having now generally described the invention, the same will be more readily understood through reference to the following examples which are provided by way of illustration and are not intended to be limiting of the present invention, unless specified.

### **Example 1**

The MONN01 cDNA library is a normalized library generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting

at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON001 cDNA library is generated from maize (B73, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) immature tassels at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in a greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue from the maize plant is collected at the V6 stage. At that stage the tassel is an immature tassel of about 2-3 cm in length. The tassels are removed and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON003 library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) roots at the V6 developmental stage. Seeds are planted at a depth of approximately 3 cm in soil into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth, the seedlings are transplanted into 10 inch pots containing the Metro 200 growing medium. Plants are watered daily before transplantation and approximately 3 times a week after transplantation. Peters 15-16-17 fertilizer is applied approximately three times per week after transplanting at a concentration of 150 ppm N. Two to three times during the life time of the plant from transplanting to flowering a total of approximately 900 mg Fe is added to each pot. Maize plants are grown in the green house in approximately 15hr day/9hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6 leaf development stage. The root system is cut from maize plant and washed with water to free it from the soil. The tissue is then immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON004 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is

approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON005 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) root tissue at the V6 development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The root system is cut from the mature maize plant and washed with water to free it from the soil. The tissue is immediately frozen in liquid nitrogen and the harvested tissue is then stored at -80°C until RNA preparation.

The SATMON006 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) total leaf tissue at the V6 plant development

stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older more juvenile leaves, which are in a basal position, as well as the younger more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON007 cDNA library is generated from the primary root tissue of 5 day old maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark until germination (one day). After germination, the trays, along with the moist paper, are moved to a greenhouse where the maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles for approximately 5 days. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. The primary root tissue is collected when the seedlings are 5 days old. At this stage, the primary root (radicle) is



pushed through the coleorhiza which itself is pushed through the seed coat. The primary root, which is about 2-3 cm long, is cut and immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON008 cDNA library is generated from the primary shoot (coleoptile 2-3 cm) of maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings which are approximately 5 days old. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark until germination (one day). Then the trays containing the seeds are moved to a greenhouse at 15hr daytime/9 hr nighttime cycles and grown until they are 5 days post germination. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Tissue is collected when the seedlings are 5 days old. At this stage, the primary shoot (coleoptile) is pushed through the seed coat and is about 2-3 cm long. The coleoptile is dissected away from the rest of the seedling, immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON009 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaves at the 8 leaf stage (V8 plant development stage). Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is 80°F and the nighttime temperature is

70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 8-leaf development stage. The older more juvenile leaves, which are in a basal position, as well as the younger more adult leaves, which are more apical, are cut at the base of the leaves. The leaves are then pooled and then immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON010 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) root tissue at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is 80°F and the nighttime temperature is 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the V8 development stage. The root system is cut from this mature maize plant and washed with water to free it from the soil. The tissue is immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON011 cDNA library is generated from undeveloped maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaf at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium.

After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The second youngest leaf which is at the base of the apical leaf of V6 stage maize plant is cut at the base and immediately transferred to liquid nitrogen containers in which the leaf is crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON012 cDNA library is generated from 2 day post germination maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark until germination (one day). Then the trays containing the seeds are moved to the greenhouse and grown at 15hr daytime/9 hr nighttime cycles until 2 days post germination. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Tissue is collected when the seedlings are 2 days old. At the two day stage, the coleorhiza is pushed through the seed coat and the primary root (the radicle) is pierced the coleorhiza but is barely visible. Also, at this two day stage, the coleoptile is just emerging from the seed coat. The 2 days post germination seedlings are then immersed in liquid nitrogen and crushed. The harvested tissue is stored at -80°C until preparation of total RNA.

The SATMON013 cDNA library is generated from apical maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) meristem founder at the V4 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Prior to tissue collection, the plant is at the 4 leaf stage. The lead at the apex of the V4 stage maize plant is referred to as the meristem founder. This apical meristem founder is cut, immediately frozen in liquid nitrogen and crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON014 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) endosperm fourteen days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9

hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the maize plant ear shoots are ready for fertilization. At this stage, the ear shoots are enclosed in a paper bag before silk emergence to withhold the pollen. The ear shoots are pollinated and 14 days after pollination, the ears are pulled out and then the kernels are plucked out of the ears. Each kernel is then dissected into the embryo and the endosperm and the aleurone layer is removed. After dissection, the endosperms are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON016 library is a maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) sheath library collected at the V8 developmental stage. Seeds are planted in a depth of approximately 3 cm in solid into 2-3 inch pots containing Metro growing medium. After 2-3 weeks growth, they are transplanted into 10" pots containing the same. Plants are watered daily before transplantation and approximately the times a week after transplantation. Peters 15-16-17 fertilizer is applied approximately three times per week after transplanting, at a strength of 150 ppm N. Two to three times during the life time of the plant from transplanting to flowering, a total of approximately 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15hr day/9hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. When the maize plants are at the V8 stage the 5<sup>th</sup> and 6<sup>th</sup> leaves from the bottom exhibit fully developed leaf blades. At the base of these leaves, the ligule is differentiated and the leaf blade is joined to the sheath. The sheath is dissected away from the

base of the leaf then the sheath is frozen in liquid nitrogen and crushed. The tissue is then stored at -80°C until RNA preparation.

The SATMON017 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) embryo seventeen days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth the seeds are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the ear shoots of maize plant, which are ready for fertilization, are enclosed in a paper bag before silk emergence to withhold the pollen. The ear shoots are fertilized and 21 days after pollination, the ears are pulled out and the kernels are plucked out of the ears. Each kernel is then dissected into the embryo and the endosperm and the aleurone layer is removed. After dissection, the embryos are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON019 (Lib3054) cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) culm (stem) at the V8 developmental stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing

medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. When the maize plant is at the V8 stage, the 5th and 6th leaves from the bottom have fully developed leaf blades. The region between the nodes of the 5th and the sixth leaves from the bottom is the region of the stem that is collected. The leaves are pulled out and the sheath is also torn away from the stem. This stem tissue is completely free of any leaf and sheath tissue. The stem tissue is then frozen in liquid nitrogen and stored at -80°C until RNA preparation.

The SATMON020 cDNA library is from a maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) Hill Type II-Initiated Callus. Petri plates containing approximately 25 ml of Type II initiation media are prepared. This medium contains N6 salts and vitamins, 3% sucrose, 2.3 g/liter proline 0.1 g/liter enzymatic casein hydrolysate, 2mg/liter 2,4 – dichloro phenoxy-acetic acid (2,4, D), 15.3 mg/liter AgNO<sub>3</sub> and 0.8% bacto agar and is adjusted to pH 6.0 before autoclaving. At 9-11 days after pollination, an ear with immature embryos measuring approximately 1-2 mm in length is chosen. The husks and silks are removed and then the ear is broken into halves and placed in an autoclaved solution of Clorox/TWEEN 20 sterilizing solution. Then the ear is rinsed with deionized water. Then each embryo is extracted from the kernel. Intact embryos are placed in contact with the medium, scutellar side up). Multiple embryos are plated on each plate and the plates are incubated in the dark at 25°C. Type II

calluses are friable, can be subcultured with a spatula, frequently regenerate via somatic embryogenesis and are relatively undifferentiated. As seen in the microscope, the Tape II calluses show color ranging from translucent to light yellow and heterogeneity on with respect to embryoid structure as well as stage of embryoid development. Once Type II callus are formed, the calluses is transferred to type II callus maintenance medium without  $\text{AgNO}_3$ . Every 7-10 days, the callus is subcultured. About 4 weeks after embryo isolation the callus is removed from the plates and then frozen in liquid nitrogen. The harvested tissue is stored at  $-80^\circ\text{C}$  until RNA preparation.

The SATMON021 cDNA library is generated from the immature maize (DK604, Dekalb Genetics, Dekalb Illinois, U.S.A.) tassel at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately  $80^\circ\text{F}$  and the nighttime temperature is approximately  $70^\circ\text{F}$ . Supplemental lighting is provided by 1000 W sodium vapor lamps. As the maize plant enters the V8 stage, tassels which are 15-20 cm in length are collected and frozen in liquid nitrogen. The harvested tissue is stored at  $-80^\circ\text{C}$  until RNA preparation.

The SATMON022 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) ear (growing silks) at the V8 plant development stage. Seeds are planted



at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. *Zea mays* plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the plant is in the V8 stage. At this stage, some immature ear shoots are visible. The immature ear shoots (approximately 1 cm in length) are pulled out, frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON23 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) ear (growing silk) at the V8 development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. When the tissue is harvested at the V8 stage, the length of the ear that is harvested is about 10-15 cm and the silks are just exposed (approximately 1 inch).

The ear along with the silks is frozen in liquid nitrogen and then the tissue is stored at -80°C until RNA preparation.

The SATMON024 cDNA library is generated from the immature maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) tassel at the V9 development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. As a maize plant enters the V9 stage, the tassel is rapidly developing and a 37 cm tassel along with the glume, anthers and pollen is collected and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The SATMON025 cDNA library is from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) Hill Type II-Regenerated Callus. Type II callus is grown in initiation media as described for SATMON020 and then the embryoids on the surface of the Type II callus are allowed to mature and germinate. The 1-2 gm fresh weight of the soft friable type callus containing numerous embryoids are transferred to 100 x 15 mm petri plates containing 25 ml of regeneration media. Regeneration media consists of Murashige and Skoog (MS) basal salts, modified White's vitamins (0.2 g/liter glycine and 0.5 g/liter myo-inositol and 0.8% bacto agar (6SMS0D)). The plates are then placed in the dark after covering with parafilm. After 1 week,

the plates are moved to a lighted growth chamber with 16 hr light and 8 hr dark photoperiod.

Three weeks after plating the Type II callus to 6SMS0D, the callus exhibit shoot formation. The callus and the shoots are transferred to fresh 6SMS0D plates for another 2 weeks. The callus and the shoots are then transferred to petri plates with reduced sucrose (3SMS0D). Upon distinct formation of a root and shoot, the newly developed green plants are then removed out with a spatula and frozen in liquid nitrogen containers. The harvested tissue is then stored at  $-80^{\circ}\text{C}$  until RNA preparation.

The SATMON026 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) juvenile/adult shift leaves at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately  $80^{\circ}\text{F}$  and the nighttime temperature is approximately  $70^{\circ}\text{F}$ . Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plants are at the 8-leaf development stage. Leaves are founded sequentially around the meristem over weeks of time and the older, more juvenile leaves arise earlier and in a more basal position than the younger, more adult leaves, which are in a more apical position. In a V8 plant, some leaves which are in the middle portion of the plant exhibit characteristics of both juvenile as well as adult leaves. They exhibit a yellowing color

but also exhibit, in part, a green color. These leaves are termed juvenile/adult shift leaves. The juvenile/adult shift leaves (the 4th, 5th leaves from the bottom) are cut at the base, pooled and transferred to liquid nitrogen in which they are then crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON027 cDNA library is generated from 6 day maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaves. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the Metro 200 growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. *Zea mays* plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Prior to tissue collection, when the plant is at the 8-leaf stage, water is held back for six days. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical, are all cut at the base of the leaves. All the leaves exhibit significant wilting. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are then crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON028 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) roots at the V8 developmental stage that are subject to six days water

stress. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the Metro 200 growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Prior to tissue collection, when the plant is at the 8-leaf stage, water is held back for six days. The root system is cut, shaken and washed to remove soil. Root tissue is then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are then crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON029 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings at the etiolated stage. Seeds are planted on a moist filter paper on a covered tray that is kept in the dark for 4 days at approximately 70°F. Tissue is collected when the seedlings are 4 days old. By 4 days, the primary root has penetrated the coleorhiza and is about 4-5 cm and the secondary lateral roots have also made their appearance. The coleoptile has also pushed through the seed coat and is about 4-5 cm long. The seedlings are frozen in liquid nitrogen and crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON030 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) root tissue at the V4 plant development stage. Seeds are planted at a

depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth, they are transplanted into 10 inch pots containing the same. Plants are watered daily before transplantation and approximately 3 times a week after transplantation. Peters 15-16-17 fertilizer is applied approximately three times per week after transplanting, at a strength of 150 ppm N. Two to three times during the life time of the plant, from transplanting to flowering, a total of approximately 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15hr day/9hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 sodium vapor lamps. Tissue is collected when the maize plant is at the 4 leaf development stage. The root system is cut from the mature maize plant and washed with water to free it from the soil. The tissue is then immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON031 cDNA library is generated from the maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) leaf tissue at the V4 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is 80°F and the nighttime temperature is 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected

when the maize plant is at the 4-leaf development stage. The third leaf from the bottom is cut at the base and immediately frozen in liquid nitrogen and crushed. The tissue is immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON033 cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) embryo tissue 13 days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the ear shoots of the maize plant, which are ready for fertilization, are enclosed in a paper bag before silk emergent to withhold the pollen. The ear shoots are pollinated and 13 days after pollination, the ears are pulled out and then the kernels are plucked cut of the ears. Each kernel is then dissected into the embryo and the endosperm and the aleurone layer is removed. After dissection, the embryos are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The SATMON034 cDNA library is generated from cold stressed maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) seedlings. Seeds are planted on a moist filter paper on a covered tray that is kept on at 10°C for 7 days. After 7 days, the temperature is shifted to 15°C

for one day until germination of the seed. Tissue is collected once the seedlings are 1 day old. At this point, the coleorhiza has just pushed out of the seed coat and the primary root is just making its appearance. The coleoptile has not yet pushed completely through the seed coat and is also just making its appearance. These 1 day old cold stressed seedlings are frozen in liquid nitrogen and crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMON~001 (Lib36, Lib83, Lib84) cDNA library is generated from maize leaves at the V8 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in a greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue from the maize plant is collected at the V8 stage. The older more juvenile leaves in a basal position as well as the younger more adult leaves which are more apical are all cut at the base, pooled and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SATMONN01 cDNA library is generated from maize (B73, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) normalized immature tassels at the V6 plant development stage normalized tissue. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into



10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in a greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue from the maize plant is collected at the V6 stage. At that stage the tassel is an immature tassel of about 2-3 cm in length. The tassels are removed and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The SATMONN04 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign, Illinois U.S.A.) normalized total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10

inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older, more juvenile leaves, which are in a basal position, as well as the younger, more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The SATMONN05 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) normalized root tissue at the V6 development

stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The root system is cut from the mature maize plant and washed with water to free it from the soil. The tissue is immediately frozen in liquid nitrogen and the harvested tissue is then stored at -80°C until RNA preparation. The single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The SATMONN06 cDNA library is generated from maize (B73 x Mo17, Illinois Foundation Seeds, Champaign Illinois, U.S.A.) normalized total leaf tissue at the V6 plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 6-leaf development stage. The older more juvenile leaves, which are in a basal position, as well as the younger more adult leaves, which are more apical are cut at the base of the leaves. The leaves are then pooled and immediately transferred to liquid nitrogen containers in which the pooled leaves are crushed. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-

hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The CMZ029 (SATMON036) cDNA library is generated from maize (DK604, Dekalb Genetics, Dekalb, Illinois U.S.A.) endosperm 22 days after pollination. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the green house in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. After the V10 stage, the ear shoots of the maize plant, which are ready for fertilization, are enclosed in a paper bag before silk emergent to withhold the pollen. The ear shoots are pollinated and 22 days after pollination, the ears are pulled out and then the kernels are plucked out of the ears. Each kernel is then dissected into the embryo and the endosperm and the alurone layer is removed. After dissection, the endosperms are immediately frozen in liquid nitrogen and then stored at -80°C until RNA preparation.

The CMz030 (Lib143) cDNA library is generated from maize seedling tissue two days post germination. Seeds are planted on a moist filter paper on a covered try that is keep in the dark until germination. The trays are then moved to the bench top at 15 hr daytime/9 hr nighttime cycles for 2 days post-germination. The day time temperature is 80°F and the

nighttime temperature is 70°F. Tissue is collected when the seedlings are 2 days old. At this stage, the colehrhiza has pushed through the seed coat and the primary root (the radicle) is just piercing the colehrhiza and is barely visible. The seedlings are placed at 42°C for 1 hour. Following the heat shock treatment, the seedlings are immersed in liquid nitrogen and crushed. The harvested tissue is stored at -80°C until RNA preparation.

The CMz031 (Lib148) cDNA library is generated from maize pollen tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants. The ear shoots, which are ready for fertilization, are enclosed in a paper bag to withhold pollen. Twenty-one days after pollination, prior to removing the ears, the paper bag is shaken to collect the mature pollen. The mature pollen is immediately frozen in liquid nitrogen containers and the pollen is crushed. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz033 (Lib189) cDNA library is generated from maize pooled leaf tissue. Samples are harvested from open pollinated plants. Tissue is collected from maize leaves at the

anthesis stage. The leaves are collected from 10-12 plants and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz034 (Lib3060) cDNA library is generated from maize mature tissue at 40 days post pollination plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from leaves located two leaves below the ear leaf. This sample represents those genes expressed during onset and early stages of leaf senescence. The leaves are pooled and immediately transferred to liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz036 (Lib3062) cDNA library is generated from maize husk tissue at the 8 week old plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during

the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from 8 week old plants. The husk is separated from the ear and immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz037 (Lib3059) cDNA library is generated from maize pooled kernal at 12-15 days after pollination plant development stage. Sample were collected from field grown material. Whole kernal from hand pollinated (control pollination) are harvested as whole ears and immediately frozen on dry ice. Kernels from 10-12 ears were pooled and ground together in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz039 (Lib3066) cDNA library is generated from maize immature anther tissue at the 7 week old immature tassel stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 7 week old immature tassel stage. At this stage, prior to anthesis, the



immature anthers are green and enclosed in the staminate spikelet. The developing anthers are dissected away from the 7 week old immature tassel and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz040 (Lib3067) cDNA library is generated from maize kernel tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants. The ear shoots, which are ready for fertilization, are enclosed in a paper bag before silk emergence to withhold pollen. Five to eight days after controlled pollination. The ears are pulled and the kernels removed. The kernels are immediately frozen in liquid nitrogen. The harvested kernels tissue is then stored at -80°C until RNA preparation. This sample represents gene expressed in early kernel development, during periods of cell division, amyloplast biogenesis and early carbon flow across the material to filial tissue.

The CMz041 (Lib3068) cDNA library is generated from maize pollen germinating silk tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are

transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants when the ear shoots are ready for fertilization at the silk emergence stage. The emerging silks are pollinated with an excess of pollen under controlled pollination conditions in the green house. Eighteen hours after pollination the silks are removed from the ears and immediately frozen in liquid nitrogen containers. This sample represents genes expressed in both pollen and silk tissue early in pollination. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz042 (Lib3069) cDNA library is generated from maize ear tissue excessively pollinated at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F.

Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ stage plants and the ear shoots which are ready for fertilization are at the silk emergence stage. The immature ears are pollinated with an excess of pollen under controlled pollination conditions. Eighteen hours post-pollination, the ears are removed and immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz044 (Lib3075) cDNA library is generated from maize microspore tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F.

Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from immature anthers from 7 week old tassels. The immature anthers are first dissected from the 7 week old tassel with a scalpel on a glass slide covered with water. The microspores (immature pollen) are released into the water and are recovered by centrifugation. The microspore suspension is immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz045 (Lib3076) cDNA library is generated from maize immature ear megaspore tissue. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing

Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from immature ear (megaspore) obtained from 7 week old plants. The immature ears are harvested from the 7 week old plants and are approximately 2.5 to 3 cm in length. The kernels are removed from the cob immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz047 (Lib3078) cDNA library is generated from maize CO<sub>2</sub> treated high-exposure shoot tissue at the V10+ plant development stage. RX601 maize seeds are sterilized for 1 minute with a 10% clorox solution. The seeds are rolled in germination paper, and germinated in 0.5 mM calcium sulfate solution for two days at 30°C. The seedlings are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium at a rate of 2-3 seedlings per pot. Twenty pots are placed into a high CO<sub>2</sub> environment (approximately 1000 ppm CO<sub>2</sub>). Twenty plants were grown under ambient greenhouse CO<sub>2</sub> (approximately 450 ppm CO<sub>2</sub>). Plants are watered daily before transplantation and three times a week after transplantation. Peters 20-20-20 fertilizer is also lightly applied. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime

temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. At ten days post planting, the shoots from both atmosphere are frozen in liquid nitrogen and lightly ground. The roots are washed in deionized water to remove the support media and the tissue is immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz048 (Lib3079) cDNA library is generated from maize basal endosperm transfer layer tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected from V10+ maize plants. The ear shoots, which are ready for fertilization, are enclosed in a paper bag prior to silk emergence, to withhold the pollen. Kernels are harvested at 12 days post-pollination and placed on wet ice for dissection. The kernels are cross sectioned laterally, dissecting just above the pedicel region, including 1-2 mm of the lower endosperm and the basal endosperm transfer region. The pedicel and lower endosperm region containing the basal endosperm transfer layer is pooled and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz049(Lib3088) cDNA library is generated from maize immature anther tissue at the 7 week old immature tassel stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime temperature is approximately 80°F and the nighttime temperature is approximately 70°F. Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is at the 7 week old immature tassel stage. At this stage, prior to anthesis, the immature anthers are green and enclosed in the staminate spikelet. The developing anthers are dissected away from the 7 week old immature tassel and immediately transferred to liquid nitrogen container. The harvested tissue is then stored at -80°C until RNA preparation.

The CMz050 (Lib3114) cDNA library is generated from maize silk tissue at the V10+ plant development stage. Seeds are planted at a depth of approximately 3 cm into 2-3 inch peat pots containing Metro 200 growing medium. After 2-3 weeks growth they are transplanted into 10 inch pots containing the same growing medium. Plants are watered daily before transplantation and three times a week after transplantation. Peters 15-16-17 fertilizer is applied three times per week after transplanting at a strength of 150 ppm N. Two to three times during the lifetime of the plant, from transplanting to flowering, a total of 900 mg Fe is added to each pot. Maize plants are grown in the greenhouse in 15 hr day/9 hr night cycles. The daytime

temperature is approximately 80°F and the nighttime temperature is approximately 70°F.

Supplemental lighting is provided by 1000 W sodium vapor lamps. Tissue is collected when the maize plant is beyond the 10-leaf development stage and the ear shoots are approximately 15-20 cm in length. The ears are pulled and silks are separated from the ears and immediately transferred to liquid nitrogen containers. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON001 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) total leaf tissue at the V4 plant development stage. Leaf tissue from 38, field grown V4 stage plants is harvested from the 4<sup>th</sup> node. Leaf tissue is removed from the plants and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON002 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue at the V4 plant development stage. Root tissue from 76, field grown V4 stage plants is harvested. The root systems is cut from the soybean plant and washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON003 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling hypocotyl axis tissue harvested 2 day post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even

moisture conditions. Tissue is collected 2 days after the start of imbibition. The 2 days after imbibition samples are separated into 3 collections after removal of any adhering seed coat. At the 2 day stage, the hypocotyl axis is emerging from the soil. A few seedlings have cracked the soil surface and exhibited slight greening of the exposed cotyledons. The seedlings are washed in water to remove soil, hypocotyl axis harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON004 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling cotyledon tissue harvested 2 day post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Tissue is collected 2 days after the start of imbibition. The 2 days after imbibition samples are separated into 3 collections after removal of any adhering seed coat. At the 2 day stage, the hypocotyl axis is emerging from the soil. A few seedlings have cracked the soil surface and exhibited slight greening of the exposed cotyledons. The seedlings are washed in water to remove soil, hypocotyl axis harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON005 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling hypocotyl axis tissue harvested 6 hour post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the



nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Tissue is collected 6 hours after the start of imbibition. The 6 hours after imbibition samples are separated into 3 collections after removal of any adhering seed coat. The 6 hours after imbibition sample is collected over the course of approximately 2 hours starting at 6 hours post imbibition. At the 6 hours after imbibition stage, not all cotyledons have become fully hydrated and germination, or radicle protrusion, has not occurred. The seedlings are washed in water to remove soil, hypocotyl axis harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON006 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling cotyledons tissue harvest 6 hour post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. Trays are placed in an environmental chamber and grown at 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Tissue is collected 6 hours after imbibition. The 6 hours after imbibition samples are separated into 3 collections after removal of any adhering seed coat. The 6 hours after imbibition sample is collected over the course of approximately 2 hours starting at 6 hours post-imbibition. At the 6 hours after imbibition, not all cotyledons have become fully hydrated and germination or radicle protrusion, have not occurred. The seedlings are washed in water to remove soil, cotyledon harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON007 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 25 and 35 days post-

flowering. Seed pods from field grown plants are harvested 25 and 35 days after flowering and the seeds extracted from the pods. Approximately 4.4g and 19.3g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON008 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue harvested from 25 and 35 days post-flowering plants. Total leaf tissue is harvested from field grown plants. Approximately 19g and 29g of leaves are harvested from the fourth node of the plant 25 and 35 days post-flowering and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON009 cDNA library is generated from soybean cutlivar C1944 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) pod and seed tissue harvested 15 days post-flowering. Pods from field grown plants are harvested 15 days post-flowering. Approximately 3g of pod tissue is harvested and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON010 cDNA library is generated from soybean cultivar C1944 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) seed tissue harvested 40 days post-flowering. Pods from field grown plants are harvested 40 days post-flowering. Pods and seeds are separated, approximately 19g of seed tissue is harvested and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON011 cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical

germ plasma) leaf tissue. Leaves are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 30g of leaves are harvested from the 4<sup>th</sup> node of each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON012 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue. Leaves from field grown plants are harvested from the fourth node 15 days post-flowering. Approximately 12g of leaves are harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON013 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root and nodule tissue. Approximately, 28g of root tissue from field grown plants is harvested 15 days post-flowering. The root system is cut from the soybean plant, washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON014 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 25 and 35 days after flowering. Seed pods from field grown plants are harvested 15 days after flowering and the seeds extracted from the pods. Approximately 5g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON015 cDNA is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 45 and 55 days post-flowering. Seed pods from field grown plants are harvested 45 and 55 days after flowering and the seeds extracted from the pods. Approximately 19g and 31g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON016 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue. Approximately, 61g and 38g of root tissue from field grown plants is harvested 25 and 35 days post-flowering is harvested. The root system is cut from the soybean plant and washed with water to free it from the soil. The tissue is placed in 14ml polystyrene tubes and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON017 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue. Approximately 28g of root tissue from field grown plants is harvested 45 and 55 days post-flowering. The root system is cut from the soybean plant, washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON018 cDNA is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue harvested 45 and 55 days post-flowering. Leaves from field grown plants are harvested 45 and 55 days after flowering from the fourth node. Approximately 27g and 33g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON019 cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical germ plasma) root tissue. Roots are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 50g and 56g of roots are harvested from each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON020 cDNA is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 65 and 75 days post-flowering. Seed pods from field grown plants are harvested 45 and 55 days after flowering and the seeds extracted from the pods. Approximately 14g and 31g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON021 cDNA library is generated from Soybean Cyst Nematode-resistant soybean cultivar Hartwig (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) root tissue. Plants are grown in tissue culture at room temperature. At approximately 6 weeks post-germination, the plants are exposed to sterilized Soybean Cyst Nematode eggs. Infection is then allowed to progress for 10 days. After the 10 day infection process, the tissue is harvested. Agar from the culture medium and nematodes are removed and the root tissue is immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON022 (Lib3030) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) partially opened flower tissue.

Partially to fully opened flower tissue is harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. A total of 3g of flower tissue is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON023 cDNA library is generated from soybean genotype BW211S Null (Tohoku University, Morioka, Japan) seed tissue harvested 15 and 40 days post-flowering. Seed pods from field grown plants are harvested 15 and 40 days post-flowering and the seeds extracted from the pods. Approximately 0.7g and 14.2g of seeds are harvested from the respective seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON024 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) internode-2 tissue harvested 18 days post-imbibition. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium. The plants are grown in a greenhouse for 18 days after the start of imbibition at ambient temperature. Soil is checked and watered daily to maintain even moisture conditions. Stem tissue is harvested 18 days after the start of imbibition. The samples are divided into hypocotyl and internodes 1 through 5. The fifth internode contains some leaf bud material. Approximately 3 g of each sample is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON025 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) leaf tissue harvested 65 days post-flowering. Leaves are harvested from the fourth node of field grown plants 65 days post-flowering.

Approximately 18.4g of leaf tissue is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

SOYMON026 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) root tissue harvested 65 and 75 days post-flowering. Approximately 27g and 40g of root tissue from field grown plants is harvested 65 and 75 days post-flowering. The root system is cut from the soybean plant, washed with water to free it from the soil and immediately frozen in dry-ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON027 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed tissue harvested 25 days post-flowering. Seed pods from field grown plants are harvested 25 days post-flowering and the seeds extracted from the pods. Approximately 17g of seeds are harvested from the seed pods and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON028 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought-stressed root tissue. The plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of development, water is withheld from half of the plant collection (drought stressed population). After 3 days, half of the plants from the drought stressed condition and half of the plants from the control population are harvested. After another 3 days (6 days post drought induction) the remaining plants are

harvested. A total of 27g and 40g of root tissue is harvested and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON029 cDNA library is generated from Soybean Cyst Nematode-resistant soybean cultivar PI07354 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) root tissue. Late fall to early winter greenhouse grown plants are exposed to Soybean Cyst Nematode eggs. At 10 days post-infection, the plants are uprooted, rinsed briefly and the roots frozen in liquid nitrogen. Approximately 20 grams of root tissue is harvested from the infected plants. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON030 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) flower bud tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Flower buds are removed from the plant at the pedicel. A total of 100mg of flower buds are harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON031 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) carpel and stamen tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Flower buds are



removed from the plant at the pedicel. Flowers are dissected to separate petals, sepals and reproductive structures (carpels and stamens). A total of 300mg of carpel and stamen tissue are harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON032 cDNA library is prepared from the Asgrow cultivar A4922 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) rehydrated dry soybean seed meristem tissue. Surface sterilized seeds are germinated in liquid media for 24 hours. The seed axis is then excised from the barely germinating seed, placed on tissue culture media and incubated overnight at 20°C in the dark. The supportive tissue is removed from the explant prior to harvest. Approximately 570mg of tissue is harvested and frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON033 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) heat-shocked seedling tissue without cotyledons. Seeds are imbibed and germinated in vermiculite for 2 days under constant illumination. After 48 hours, the seedlings are transferred to an incubator set at 40°C under constant illumination. After 30, 60 and 180 minutes seedlings are harvested and dissected. A portion of the seedling consisting of the root, hypocotyl and apical hook is frozen in liquid nitrogen and stored at -80°C. The seedlings after 2 days of imbibition are beginning to emerge from the vermiculite surface. The apical hooks are dark green in appearance. Total RNA and poly A<sup>+</sup> RNA is prepared from equal amounts of pooled tissue.

The SOYMON034 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) cold-shocked seedling tissue without cotyledons. Seeds are imbibed and germinated in vermiculite for 2 days under constant

illumination. After 48 hours, the seedlings are transferred to a cold room set at 5°C under constant illumination. After 30, 60 and 180 minutes seedlings are harvested and dissected. A portion of the seedling consisting of the root, hypocotyl and apical hook is frozen in liquid nitrogen and stored at -80°C. The seedlings after 2 days of imbibition are beginning to emerge from the vermiculite surface. The apical hooks are dark green in appearance.

The SOYMON035 cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seed coat tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Seeds are harvested from mid to nearly full maturation (seed coats are not yellowing). The entire embryo proper is removed from the seed coat sample and the seed coat tissue are harvested and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON036 cDNA library is generated from soybean cultivars PI171451, PI227687 and PI229358 (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) insect challenged leaves. Plants from each of the three cultivars are grown in screenhouse conditions. The screenhouse is divided in half and one half of the screenhouse is infested with soybean looper and the other half infested with velvetbean caterpillar. A single leaf is taken from each of the representative plants at 3 different time points, 11 days after infestation, 2 weeks after infestation and 5 weeks after infestation and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation. Total RNA and poly A+ RNA is

isolated from pooled tissue consisting of equal quantities of all 18 samples (3 genotypes X 3 sample times X 2 insect genotypes).

The SOYMON037 cDNA library is generated from soybean cultivar A3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) etiolated axis and radical tissue. Seeds are planted in moist vermiculite, wrapped and kept at room temperature in complete darkness until harvest. Etiolated axis and hypocotyl tissue is harvested at 2, 3 and 4 days post-planting. A total of 1 gram of each tissue type is harvested at 2, 3 and 4 days after planting and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The SOYMON038 cDNA library is generated from soybean variety Asgrow A3237 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) rehydrated dry seeds. Explants are prepared for transformation after germination of surface-sterilized seeds on solid tissue media. After 6 days, at 28°C and 18 hours of light per day, the germinated seeds are cold shocked at 4°C for 24 hours. Meristemic tissue and part of the hypocotyl is removed and cotyledon excised. The prepared explant is then wounded for *Agrobacterium* infection. The 2 grams of harvested tissue is frozen in liquid nitrogen and stored at -80°C until RNA preparation.

The Soy51 (LIB3027) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The

dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The Soy52 (LIB3028) cDNA library is generated from normalized flower DNA. Single stranded DNA representing approximately  $1 \times 10^6$  colony forming units of SOYMON022 harvested tissue is used as the starting material for normalization. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

The Soy53 (LIB3039) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) seedling shoot apical meristem tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Apical tissue is harvested from seedling shoot meristem tissue, 7-8 days after the start of imbibition. The apex of each seedling is dissected to include the fifth node to the apical meristem. The fifth node corresponds to the third trifoliate leaf in the very early stages of development. Stipules completely envelop the leaf primordia at this time. A total of 200mg of apical tissue is harvested

and immediately frozen in liquid nitrogen. The harvested tissue is then stored at -80°C until RNA preparation.

The Soy54 (LIB3040) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) heart to torpedo stage embryo tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Seeds are collected and embryos removed from surrounding endosperm and maternal tissues. Embryos from globular to young torpedo stages (by corresponding analogy to *Arabidopsis*) are collected with a bias towards the middle of this spectrum. Embryos which are beginning to show asymmetric development of cotyledons are considered the upper developmental boundary for the collection and are excluded. A total of 12 mg embryo tissue is frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

Soy55 (LIB3049) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) young seed tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Seeds are collected from very young pods (5 to 15 days after flowering). A total of 100mg of seeds are harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

Soy56 (LIB3029) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are not converted to double stranded form and represent a non-normalized seed pool for comparison to Soy51 cDNA libraries.

The Soy58 (LIB3050) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed root tissue subtracted from control root tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days root tissue from both drought stressed and control (watered regularly) plants are collected and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that

described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

The Soy59 (LIB3051) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) endosperm tissue. Seeds are germinated on paper towels under laboratory ambient light conditions. At 8, 10 and 14 hours after imbibition, the seed coats are harvested. The endosperm consists of a very thin layer of tissue affixed to the inside of the seed coat. The seed coat and endosperm are frozen immediately after harvest in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The Soy60 (LIB3072) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed seed plus pod subtracted from control seed plus pod tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 26°C and the nighttime temperature 21°C and 70% relative humidity. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days seeds and pods from both drought stressed and

control (watered regularly) plants are collected from the fifth and sixth node and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

The Soy61 (LIB3073) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid treated seedling subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St. Louis, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. After 18hours, 24hours and 48 hours post



treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). For this library's construction, the eighth fraction of the cDNA size fractionation step was used for ligation.

The Soy62 (LIB3074) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid treated seedling subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St.

Loius, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. After 18 hours, 24 hours and 48 hours post treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). For this library's construction, the ninth fraction of the cDNA size fractionation step was used for ligation.

The Soy65 (LIB3107) 07cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought-stressed abscission zone tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr

nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature 24°C. Soil is checked and watered daily to maintain even moisture conditions. Plants are irrigated with 15-16-17 Peter's Mix. At the R3 stage of development, drought is imposed by withholding water. At 3, 4, 5 and 6 days, tissue is harvested and wilting is not obvious until the fourth day. Abscission layers from reproductive organs are harvested by cutting less than one millimeter proximal and distal to the layer and immediately frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

The Soy66 (LIB3109) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) non-drought stressed abscission zone tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Plants are irrigated with 15-16-17 Peter's Mix. At 3, 4, 5 and 6 days, control abscission layer tissue is harvested. Abscission layers from reproductive organs are harvested by cutting less than one millimeter proximal and distal to the layer and immediately frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation.

Soy67 (LIB3065) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar

ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. Captured hybrids are eluted with water.

Soy68 (LIB3052) cDNA library is prepared from equal amounts tissue harvested from SOYMON007, SOYMON015 and SOYMON020 prepared tissue. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. Captured hybrids are eluted with water.

Soy69 (LIB3053) cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical germ plasma) normalized leaf tissue. Leaves are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 30g of leaves are harvested from the 4<sup>th</sup> node of each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation. Single stranded and double stranded DNA representing approximately  $1 \times 10^6$  colony forming units are isolated using standard protocols. RNA, complementary to the single stranded DNA, is synthesized using the double stranded DNA as a template. Biotinylated dATP is incorporated into the RNA during the

synthesis reaction. The single stranded DNA is mixed with the biotinylated RNA in a 1:10 molar ratio) and allowed to hybridize. DNA-RNA hybrids are captured on Dynabeads M280 streptavidin (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The dynabeads with captured hybrids are collected with a magnet. The non-hybridized single stranded molecules remaining after hybrid capture are converted to double stranded form and represent the primary normalized library.

Soy70 (LIB3055) cDNA library is generated from soybean cultivars Cristalina (USDA Soybean Germplasm Collection, Urbana, Illinois U.S.A.) and FT108 (Monsoy, Brazil) (tropical germ plasma) leaf tissue. Leaves are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 30g of leaves are harvested from the 4<sup>th</sup> node of each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

Soy71 (LIB3056) cDNA library is generated from soybean cultivars Cristalina and FT108 (tropical germ plasma) root tissue. Roots are harvested from plants grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 29°C and the nighttime temperature approximately 24°C. Soil is checked and watered daily to maintain even moisture conditions. Approximately 50g and 56g of roots are harvested from each of the Cristalina and FT108 cultivars and immediately frozen in dry ice. The harvested tissue is then stored at -80°C until RNA preparation.

Soy72 (LIB3093) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed leaf control tissue. Seeds

are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under 12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 26°C and the nighttime temperature 21°C and 70% relative humidity. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days seeds and pods from both drought stressed and control (watered regularly) plants are collected from the fifth and sixth node and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

Soy73 (LIB3093) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) drought stressed leaf subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in an environmental chamber under

12hr daytime/12hr nighttime cycles. The daytime temperature is approximately 26°C and the nighttime temperature 21°C and 70% relative humidity. Soil is checked and watered daily to maintain even moisture conditions. At the R3 stage of the plant drought is induced by withholding water. After 3 and 6 days seeds and pods from both drought stressed and control (watered regularly) plants are collected from the fifth and sixth node and frozen in dry-ice. The harvested tissue is stored at -80°C until RNA preparation. For subtraction, target cDNA is made from the drought stressed tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.).

The Soy76 (Lib3106) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid and arachidonic treated seedling subtracted from control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the

plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St. Louis, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. Arachidonic treated seedlings are sprayed with 1m/ml arachidonic acid in 0.1% Tween-20. After 18hours, 24hours and 48 hours post treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. The RNA from the arachidonic treated seedlings is isolated separately. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). Fraction 10 of the size fractionated cDNA is ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.) in order to capture some of the smaller transcripts characteristic of antifungal proteins.



Soy77 (LIB3108) cDNA library is generated from soybean cultivar Asgrow 3244 (Asgrow Seed Company, Des Moines, Iowa U.S.A.) jasmonic acid control tissue. Seeds are planted at a depth of approximately 2cm into 2-3 inch peat pots containing Metromix 350 medium and the plants are grown in a greenhouse. The daytime temperature is approximately 29.4°C and the nighttime temperature 20°C. Soil is checked and watered daily to maintain even moisture conditions. At 9 days post planting, the plantlets are sprayed with either control buffer of 0.1% Tween-20 or jasmonic acid (Sigma J-2500, Sigma, St. Louis, Missouri U.S.A.) at 1 mg/ml in 0.1% Tween-20. Plants are sprayed until runoff and the soil and the stem is soaked with the spraying solution. At 18 hours post application of jasmonic acid, the soybean plantlets appear growth retarded. Arachidonic treated seedlings are sprayed with 1m/ml arachidonic acid in 0.1% Tween-20. After 18 hours, 24 hours and 48 hours post treatment, the cotyledons are removed and the remaining leaf and stem tissue above the soil is harvested and frozen in liquid nitrogen. The harvested tissue is stored at -80°C until RNA preparation. To make RNA, the three sample timepoints were combined and ground. The RNA from the arachidonic treated seedlings is isolated separately. For subtraction, target cDNA is made from the jasmonic acid treated tissue total RNA using the SMART cDNA synthesis system from Clontech (Clontech Laboratories, Palo Alto, California U.S.A.). Driver first strand cDNA is covalently linked to Dynabeads following a protocol similar to that described in the Dynal literature (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.). The target cDNA is then heat denatured and the second strand trapped using Dynabeads oligo-dT. The target second strand cDNA is then hybridized to the driver cDNA in 400 µl 2X SSPE for two rounds of hybridization at 65°C and 20 hours. After each hybridization, the hybridization solution is removed from the system and the hybridized target cDNA removed from the driver by heat denaturation in water. After

hybridization, the remaining cDNA is trapped with Dynabeads oligo-dT. The trapped cDNA is then amplified as in previous PCR based libraries and the resulting cDNA ligated into the pSPORT vector (Invitrogen, Carlsbad California U.S.A.). Fraction 10 of the size fractionated cDNA is ligated into the pSPORT vector in order to capture some of the smaller transcripts characteristic of antifungal proteins.

The stored RNA is purified using Trizol reagent from Life Technologies (Gibco BRL, Life Technologies, Gaithersburg, Maryland U.S.A.), essentially as recommended by the manufacturer. Poly A<sup>+</sup> RNA (mRNA) is purified using magnetic oligo dT beads essentially as recommended by the manufacturer (Dynabeads, Dynal Corporation, Lake Success, New York U.S.A.).

Construction of plant cDNA libraries is well-known in the art and a number of cloning strategies exist. A number of cDNA library construction kits are commercially available. The Superscript<sup>TM</sup> Plasmid System for cDNA synthesis and Plasmid Cloning (Gibco BRL, Life Technologies, Gaithersburg, Maryland U.S.A.) is used, following the conditions suggested by the manufacturer.

Normalized libraries are made using essentially the Soares procedure (Soares *et al.*, *Proc. Natl. Acad. Sci. (U.S.A.)* 91:9228-9232 (1994), the entirety of which is herein incorporated by reference). This approach is designed to reduce the initial 10,000-fold variation in individual cDNA frequencies to achieve abundances within one order of magnitude while maintaining the overall sequence complexity of the library. In the normalization process, the prevalence of high-abundance cDNA clones decreases dramatically, clones with mid-level abundance are relatively unaffected and clones for rare transcripts are effectively increased in abundance.

## Example 2

The cDNA libraries are plated on LB agar containing the appropriate antibiotics for selection and incubated at 37° for a sufficient time to allow the growth of individual colonies. Single colonies are individually placed in each well of a 96-well microtiter plates containing LB liquid including the selective antibiotics. The plates are incubated overnight at approximately 37°C with gentle shaking to promote growth of the cultures. The plasmid DNA is isolated from each clone using Qiaprep plasmid isolation kits, using the conditions recommended by the manufacturer (Qiagen Inc., Santa Clara, California U.S.A.).

Template plasmid DNA clones are used for subsequent sequencing. For sequencing, the ABI PRISM dRhodamine Terminator Cycle Sequencing Ready Reaction Kit with AmpliTaq® DNA Polymerase, FS, is used (PE Applied Biosystems, Foster City, California U.S.A.).

## Example 3

Nucleic acid sequences that encode for the following proteins: methionine adenosyltransferase, S-adenosylmethionine decarboxylase, aspartate kinase, aspartate-semialdehyde dehydrogenase, *O*-succinylhomoserine (thiol)-lyase, cystathionine  $\beta$ -lyase, 5-methyltetrahydropteroyltriglutamate, adenosylhomocysteinase, cystathionine  $\beta$ -synthase, cystathionine  $\gamma$ -lyase and *O*-acetylhomoserine (thiol)-lyase are identified from the Monsanto EST PhytoSeq database using TBLASTN (default values)(TBLASTN compares a protein query against the six reading frames of a nucleic acid sequence). Matches found with BLAST P values equal or less than 0.001 (probability) or BLAST Score of equal or greater than 90 are classified as hits. If the program used to determine the hit is HMMSW then the score refers to HMMSW score.

In addition, the GenBank database is searched with BLASTN and BLASTX (default values) using ESTs as queries. EST that pass the hit probability threshold of  $10e^{-8}$  for the following enzymes are combined with the hits generated by using TBLASTN (described above) and classified by enzyme (see Table A below).

A cluster refers to a set of overlapping clones in the PhytoSeq database. Such an overlapping relationship among clones is designated as a “cluster” when BLAST scores from pairwise sequence comparisons of the member clones meets a predetermined minimum value or product score of 50 or more (Product Score = (BLAST SCORE x Percentage Identity)/(5 x minimum [length (Seq1), length (Seq2)]))

Since clusters are formed on the basis of single-linkage relationships, it is possible for two non-overlapping clones to be members of the same cluster if, for instance, they both overlap a third clone with at least the predetermined minimum BLAST score (stringency). A cluster ID is arbitrarily assigned to all of those clones which belong to the same cluster at a given stringency and a particular clone will belong to only one cluster at a given stringency. If a cluster contains only a single clone (a “singleton”), then the cluster ID number will be negative, with an absolute value equal to the clone ID number of its single member. Clones grouped in a cluster in most cases represent a contiguous sequence.

TABLE A\*

METHIONINE ADENOSYLTRANSFERASE ( EC 2.5.1.6 )								
Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1	-700019427	700019427H1	SATMON001	g17262	BLASTX	120	1e-11	92
2	-700074004	700074004H1	SATMON007	g1778820	BLASTN	830	1e-60	80
3	-700149718	700149718H1	SATMON007	g1778820	BLASTN	263	1e-13	80
4	-700220251	700220251H1	SATMON011	g1127582	BLASTN	243	1e-9	90
5	-700458196	700458196H1	SATMON029	g1778820	BLASTN	202	1e-22	88
6	-701172459	701172459H1	SATMONN05	g882334	BLASTN	801	1e-57	89
7	-L1436317	LIB143-062-Q1-E1-B6	LIB143	g1778820	BLASTN	341	1e-19	77
8	-L1893126	LIB189-021-Q1-E1-F4	LIB189	g1778820	BLASTN	629	1e-41	88
9	-L1894542	LIB189-033-Q1-E1-H12	LIB189	g1778820	BLASTN	564	1e-60	76
10	-L30622839	LIB3062-028-Q1-K1-C2	LIB3062	g1778820	BLASTN	625	1e-65	74
11	-L30671966	LIB3067-036-Q1-K1-D8	LIB3067	g1655576	BLASTX	118	1e-25	85
12	-L30682166	LIB3068-004-Q1-K1-D5	LIB3068	g497900	BLASTX	111	1e-29	47
13	1	LIB143-036-Q1-E1-G7	LIB143	g450548	BLASTN	1766	1e-138	88
14	1	LIB3060-013-Q1-K1-F7	LIB3060	g1778820	BLASTN	1702	1e-133	90
15	1	LIB143-013-Q1-E1-A6	LIB143	g1778820	BLASTN	1712	1e-133	88
16	1	LIB148-007-Q1-E1-B8	LIB148	g450548	BLASTN	1693	1e-132	88
17	1	LIB3079-003-Q1-K1-G1	LIB3079	g1778820	BLASTN	1236	1e-130	88
18	1	LIB3066-011-Q1-K1-E4	LIB3066	g1778820	BLASTN	1661	1e-129	86
19	1	LIB3068-008-Q1-K1-D2	LIB3068	g450548	BLASTN	1192	1e-128	88
20	1	LIB143-061-Q1-E1-E7	LIB143	g1778820	BLASTN	1638	1e-127	88
21	1	LIB189-006-Q1-E1-B2	LIB189	g1778820	BLASTN	1627	1e-126	86
22	1	LIB143-031-Q1-E1-C7	LIB143	g450548	BLASTN	1595	1e-124	86
23	1	LIB3068-011-Q1-K1-B12	LIB3068	g450548	BLASTN	1552	1e-123	87
24	1	LIB143-013-Q1-E1-G11	LIB143	g450548	BLASTN	1585	1e-123	88
25	1	LIB3062-057-Q1-K1-H9	LIB3062	g450548	BLASTN	1577	1e-122	89
26	1	LIB143-008-Q1-E1-G9	LIB143	g450548	BLASTN	1556	1e-120	89
27	1	LIB3068-058-Q1-K1-B7	LIB3068	g450548	BLASTN	1470	1e-118	86
28	1	LIB3067-048-Q1-K1-G5	LIB3067	g1778820	BLASTN	1518	1e-117	89
29	1	LIB143-049-Q1-E1-A12	LIB143	g1778820	BLASTN	1480	1e-114	91
30	1	LIB3062-010-Q1-K1-C7	LIB3062	g1778820	BLASTN	1145	1e-113	87
31	1	LIB3068-055-Q1-K1-E10	LIB3068	g450548	BLASTN	1451	1e-112	88
32	1	LIB189-025-Q1-E1-H1	LIB189	g1778820	BLASTN	1460	1e-112	90
33	1	LIB143-066-Q1-E1-F4	LIB143	g1778820	BLASTN	1309	1e-111	86
34	1	LIB143-003-Q1-E1-D10	LIB143	g1778820	BLASTN	1440	1e-111	87
35	1	LIB3062-033-Q1-K1-G1	LIB3062	g1778820	BLASTN	1449	1e-111	86
36	1	LIB3066-029-Q1-K1-H11	LIB3066	g1778820	BLASTN	1427	1e-110	88
37	1	LIB3068-016-Q1-K1-D9	LIB3068	g450548	BLASTN	1213	1e-109	87
38	1	LIB3062-027-Q1-K1-B11	LIB3062	g1778820	BLASTN	1187	1e-108	87
39	1	LIB3068-048-Q1-K1-G4	LIB3068	g1778820	BLASTN	1410	1e-108	82
40	1	LIB148-020-Q1-E1-B6	LIB148	g1778820	BLASTN	1088	1e-107	84
41	1	LIB3066-046-Q1-K1-F2	LIB3066	g1778820	BLASTN	1400	1e-107	90
42	1	700084130H1	SATMON011	g1778820	BLASTN	960	1e-106	92
43	1	700092863H1	SATMON008	g1778820	BLASTN	1388	1e-106	92
44	1	LIB3068-043-Q1-K1-A12	LIB3068	g450548	BLASTN	827	1e-105	81
45	1	700092659H1	SATMON008	g1778820	BLASTN	1359	1e-104	92
46	1	LIB143-038-Q1-E1-H11	LIB143	g450548	BLASTN	1345	1e-103	84
47	1	LIB3062-028-Q1-K1-F12	LIB3062	g1778820	BLASTN	1082	1e-102	82
48	1	700103135H1	SATMON010	g1778820	BLASTN	1306	1e-100	93

49	1	LIB3078-050-Q1-K1-D9	LIB3078	g1778820	BLASTN	1309	1e-100	86
50	1	700084823H1	SATMON011	g1778820	BLASTN	1311	1e-100	90
51	1	700201311H1	SATMON003	g1778820	BLASTN	960	1e-97	89
52	1	700265914H1	SATMON017	g1778820	BLASTN	1275	1e-97	91
53	1	700205948H1	SATMON003	g1778820	BLASTN	983	1e-96	93
54	1	LIB189-003-Q1-E1-A6	LIB189	g450548	BLASTN	1262	1e-96	86
55	1	LIB143-017-Q1-E1-B9	LIB143	g1778820	BLASTN	779	1e-95	86
56	1	700097434H1	SATMON009	g450548	BLASTN	1247	1e-95	90
57	1	700089089H1	SATMON011	g450548	BLASTN	1251	1e-95	89
58	1	700071949H1	SATMON007	g1778820	BLASTN	1256	1e-95	87
59	1	700047892H1	SATMON003	g1778820	BLASTN	1236	1e-94	91
60	1	700077246H1	SATMON007	g450548	BLASTN	1237	1e-94	89
61	1	700085681H1	SATMON011	g1778820	BLASTN	1237	1e-94	89
62	1	LIB3067-010-Q1-K1-A7	LIB3067	g1778820	BLASTN	1244	1e-94	83
63	1	700619961H1	SATMON034	g450548	BLASTN	962	1e-93	89
64	1	700087395H1	SATMON011	g450548	BLASTN	1040	1e-93	89
65	1	700240431H1	SATMON010	g1778820	BLASTN	1156	1e-93	92
66	1	700053822H1	SATMON011	g1778820	BLASTN	1224	1e-93	91
67	1	700103856H1	SATMON010	g450548	BLASTN	1224	1e-93	88
68	1	700083703H1	SATMON011	g450548	BLASTN	1229	1e-93	89
69	1	700087360H1	SATMON011	g450548	BLASTN	1234	1e-93	90
70	1	700104713H1	SATMON010	g450548	BLASTN	1188	1e-92	87
71	1	700265996H1	SATMON017	g450548	BLASTN	1214	1e-92	88
72	1	700264870H1	SATMON017	g450548	BLASTN	945	1e-91	89
73	1	700219348H1	SATMON011	g1778820	BLASTN	1203	1e-91	91
74	1	700095527H1	SATMON008	g450548	BLASTN	1153	1e-90	88
75	1	LIB3068-023-Q1-K1-G5	LIB3068	g1778820	BLASTN	1187	1e-90	89
76	1	700102547H1	SATMON010	g1778820	BLASTN	1194	1e-90	88
77	1	700332179H1	SATMON019	g450548	BLASTN	1054	1e-89	88
78	1	700405470H1	SATMON029	g450548	BLASTN	1071	1e-89	90
79	1	LIB3069-043-Q1-K1-G2	LIB3069	g450548	BLASTN	1175	1e-89	84
80	1	700451332H1	SATMON028	g450548	BLASTN	1176	1e-89	90
81	1	700028108H1	SATMON003	g450548	BLASTN	1180	1e-89	89
82	1	700076154H1	SATMON007	g1778820	BLASTN	1180	1e-89	89
83	1	700089770H1	SATMON011	g450548	BLASTN	1185	1e-89	89
84	1	700051882H1	SATMON003	g1778820	BLASTN	844	1e-88	91
85	1	700242942H1	SATMON010	g1778820	BLASTN	1163	1e-88	91
86	1	700094996H1	SATMON008	g1778820	BLASTN	1166	1e-88	88
87	1	700476239H1	SATMON025	g450548	BLASTN	1166	1e-88	89
88	1	700219530H1	SATMON011	g1778820	BLASTN	1005	1e-87	91
89	1	700049726H1	SATMON003	g1778820	BLASTN	1041	1e-87	89
90	1	700096904H1	SATMON008	g1778820	BLASTN	1157	1e-87	86
91	1	700344026H1	SATMON021	g450548	BLASTN	1158	1e-87	88
92	1	700105590H1	SATMON010	g1778820	BLASTN	1139	1e-86	89
93	1	700465283H1	SATMON025	g450548	BLASTN	1145	1e-86	89
94	1	700104341H1	SATMON010	g450548	BLASTN	1146	1e-86	87
95	1	LIB3062-021-Q1-K1-F12	LIB3062	g1778820	BLASTN	923	1e-85	78
96	1	700105246H1	SATMON010	g450548	BLASTN	930	1e-85	89
97	1	700072146H1	SATMON007	g1778820	BLASTN	1131	1e-85	91
98	1	700028553H1	SATMON003	g1778820	BLASTN	1132	1e-85	88
99	1	700451419H1	SATMON028	g450548	BLASTN	1133	1e-85	89
100	1	700103241H1	SATMON010	g450548	BLASTN	1134	1e-85	90
101	1	700612549H1	SATMON033	g450548	BLASTN	1134	1e-85	89
102	1	700050434H1	SATMON003	g450548	BLASTN	723	1e-84	89

103	1	700094764H1	SATMON008	g1778820	BLASTN	977	1e-84	86
104	1	700163030H1	SATMON013	g1778820	BLASTN	1011	1e-84	93
105	1	700466542H1	SATMON025	g450548	BLASTN	1085	1e-84	89
106	1	700047380H1	SATMON003	g450548	BLASTN	1096	1e-84	89
107	1	700456429H1	SATMON029	g450548	BLASTN	1118	1e-84	88
108	1	700096049H1	SATMON008	g1778820	BLASTN	1122	1e-84	86
109	1	700075906H1	SATMON007	g1778820	BLASTN	838	1e-83	86
110	1	LIB3067-048-Q1-K1-G3	LIB3067	g1778820	BLASTN	1045	1e-83	86
111	1	700214043H1	SATMON016	g450548	BLASTN	1103	1e-83	88
112	1	700096909H1	SATMON008	g1778820	BLASTN	1103	1e-83	90
113	1	701184635H1	SATMONN06	g1778820	BLASTN	1104	1e-83	87
114	1	700158969H1	SATMON012	g1778820	BLASTN	1107	1e-83	93
115	1	700475902H1	SATMON025	g450548	BLASTN	1112	1e-83	89
116	1	700207076H1	SATMON003	g1778820	BLASTN	1113	1e-83	90
117	1	700161901H1	SATMON012	g1778820	BLASTN	1091	1e-82	91
118	1	700452183H1	SATMON028	g450548	BLASTN	1095	1e-82	88
119	1	700452326H1	SATMON028	g450548	BLASTN	1098	1e-82	88
120	1	700106918H1	SATMON010	g450548	BLASTN	1101	1e-82	88
121	1	700220146H1	SATMON011	g450548	BLASTN	1083	1e-81	89
122	1	700221060H1	SATMON011	g450548	BLASTN	1084	1e-81	88
123	1	700243344H1	SATMON010	g1778820	BLASTN	1084	1e-81	92
124	1	700475790H1	SATMON025	g450548	BLASTN	1084	1e-81	89
125	1	700452691H1	SATMON028	g1778820	BLASTN	1086	1e-81	87
126	1	700243733H1	SATMON010	g450548	BLASTN	1086	1e-81	90
127	1	LIB3059-018-Q1-K1-D10	LIB3059	g1778820	BLASTN	830	1e-80	91
128	1	700165958H1	SATMON013	g450548	BLASTN	1070	1e-80	89
129	1	700082659H1	SATMON011	g1778820	BLASTN	1071	1e-80	88
130	1	700172557H1	SATMON013	g1778820	BLASTN	1074	1e-80	92
131	1	700221526H1	SATMON011	g450548	BLASTN	1075	1e-80	88
132	1	700456696H1	SATMON029	g450548	BLASTN	1077	1e-80	89
133	1	700430776H1	SATMONN01	g1778820	BLASTN	574	1e-79	90
134	1	LIB3062-025-Q1-K1-B11	LIB3062	g450548	BLASTN	585	1e-79	85
135	1	700077126H1	SATMON007	g1778820	BLASTN	743	1e-79	88
136	1	700160310H1	SATMON012	g1778820	BLASTN	752	1e-79	91
137	1	700030466H1	SATMON003	g1778820	BLASTN	757	1e-79	83
138	1	700210204H1	SATMON016	g1778820	BLASTN	844	1e-79	91
139	1	700094716H1	SATMON008	g1778820	BLASTN	857	1e-79	86
140	1	700157051H1	SATMON012	g1778820	BLASTN	937	1e-79	89
141	1	700241415H1	SATMON010	g450548	BLASTN	1055	1e-79	90
142	1	700160127H1	SATMON012	g450548	BLASTN	1056	1e-79	88
143	1	700205485H1	SATMON003	g1778820	BLASTN	1059	1e-79	90
144	1	700475930H1	SATMON025	g450548	BLASTN	1060	1e-79	89
145	1	700161902H1	SATMON012	g1778820	BLASTN	1060	1e-79	91
146	1	700213573H1	SATMON016	g450548	BLASTN	1061	1e-79	89
147	1	700207677H1	SATMON016	g450548	BLASTN	1061	1e-79	88
148	1	700581117H1	SATMON031	g450548	BLASTN	1065	1e-79	88
149	1	700451418H1	SATMON028	g450548	BLASTN	546	1e-78	89
150	1	700212606H1	SATMON016	g1778820	BLASTN	1044	1e-78	85
151	1	700222605H1	SATMON011	g1778820	BLASTN	1047	1e-78	86
152	1	700050542H1	SATMON003	g1778820	BLASTN	1053	1e-78	86
153	1	700450709H1	SATMON028	g450548	BLASTN	834	1e-77	88
154	1	700213368H1	SATMON016	g450548	BLASTN	861	1e-77	89
155	1	700095926H1	SATMON008	g1778820	BLASTN	1030	1e-77	86
156	1	700261686H1	SATMON017	g1778820	BLASTN	1034	1e-77	89

157	1	700235993H1	SATMON010	g450548	BLASTN	1037	1e-77	88
158	1	700466152H1	SATMON025	g450548	BLASTN	1038	1e-77	89
159	1	700266410H1	SATMON017	g1778820	BLASTN	1041	1e-77	85
160	1	700216991H1	SATMON016	g1778820	BLASTN	491	1e-76	92
161	1	700464957H1	SATMON025	g450548	BLASTN	607	1e-76	88
162	1	700160463H1	SATMON012	g1778820	BLASTN	1018	1e-76	91
163	1	700158224H1	SATMON012	g1778820	BLASTN	1018	1e-76	92
164	1	700469971H1	SATMON025	g1778820	BLASTN	1020	1e-76	86
165	1	700222731H1	SATMON011	g450548	BLASTN	1022	1e-76	88
166	1	700087434H1	SATMON011	g1778820	BLASTN	1023	1e-76	86
167	1	700094482H1	SATMON008	g1778820	BLASTN	1023	1e-76	86
168	1	700235976H1	SATMON010	g450548	BLASTN	1028	1e-76	89
169	1	700088266H1	SATMON011	g1778820	BLASTN	983	1e-75	92
170	1	700075882H1	SATMON007	g1778820	BLASTN	1009	1e-75	83
171	1	700243324H1	SATMON010	g1778820	BLASTN	1012	1e-75	86
172	1	700159625H2	SATMON012	g450548	BLASTN	1015	1e-75	90
173	1	700204422H1	SATMON003	g1778820	BLASTN	1015	1e-75	88
174	1	700048228H1	SATMON003	g1778820	BLASTN	741	1e-74	85
175	1	700072474H1	SATMON007	g1778820	BLASTN	799	1e-74	85
176	1	700477790H1	SATMON025	g450548	BLASTN	960	1e-74	89
177	1	LIB3059-022-Q1-K1-A9	LIB3059	g1778820	BLASTN	995	1e-74	86
178	1	700241633H1	SATMON010	g450548	BLASTN	997	1e-74	89
179	1	700093978H1	SATMON008	g1778820	BLASTN	999	1e-74	89
180	1	700238569H1	SATMON010	g450548	BLASTN	844	1e-73	88
181	1	700455160H1	SATMON029	g450548	BLASTN	849	1e-73	86
182	1	700262745H1	SATMON017	g1778820	BLASTN	984	1e-73	85
183	1	700405029H1	SATMON027	g1778820	BLASTN	990	1e-73	90
184	1	700195230H1	SATMON014	g450548	BLASTN	991	1e-73	88
185	1	700264753H1	SATMON017	g1778820	BLASTN	993	1e-73	90
186	1	700050142H1	SATMON003	g1778820	BLASTN	433	1e-72	86
187	1	700452677H1	SATMON028	g450548	BLASTN	919	1e-72	87
188	1	700168870H1	SATMON013	g1778820	BLASTN	971	1e-72	90
189	1	700023193H1	SATMON003	g450548	BLASTN	972	1e-72	88
190	1	700086582H1	SATMON011	g1778820	BLASTN	980	1e-72	84
191	1	700457710H1	SATMON029	g450548	BLASTN	982	1e-72	87
192	1	700477601H1	SATMON025	g450548	BLASTN	635	1e-71	88
193	1	700377730H1	SATMON019	g1778820	BLASTN	691	1e-71	87
194	1	700169782H1	SATMON013	g1778820	BLASTN	960	1e-71	91
195	1	700461113H1	SATMON033	g1778820	BLASTN	962	1e-71	85
196	1	700258669H1	SATMON017	g1778820	BLASTN	963	1e-71	90
197	1	700454253H1	SATMON029	g1778820	BLASTN	964	1e-71	85
198	1	700092376H1	SATMON008	g1778820	BLASTN	780	1e-70	83
199	1	700242885H1	SATMON010	g960356	BLASTN	946	1e-70	86
200	1	700094333H1	SATMON008	g1778820	BLASTN	947	1e-70	86
201	1	700575124H1	SATMON030	g450548	BLASTN	948	1e-70	82
202	1	700072544H1	SATMON007	g1778820	BLASTN	949	1e-70	86
203	1	LIB143-046-Q1-E1-B5	LIB143	g450548	BLASTN	951	1e-70	87
204	1	700096534H1	SATMON008	g1778820	BLASTN	956	1e-70	85
205	1	700262906H1	SATMON017	g1778820	BLASTN	937	1e-69	87
206	1	700094478H1	SATMON008	g1778820	BLASTN	939	1e-69	90
207	1	700172985H1	SATMON013	g450548	BLASTN	939	1e-69	88
208	1	700097938H1	SATMON009	g1778820	BLASTN	941	1e-69	85
209	1	LIB3069-006-Q1-K1-D5	LIB3069	g450548	BLASTN	943	1e-69	88
210	1	700093125H1	SATMON008	g1778820	BLASTN	945	1e-69	88



211	1	700424156H1	SATMONN01	g450548	BLASTN	752	1e-68	85
212	1	700405486H1	SATMON029	g450548	BLASTN	864	1e-68	91
213	1	LIB189-026-Q1-E1-H12	LIB189	g1778820	BLASTN	879	1e-68	87
214	1	700106172H1	SATMON010	g1778820	BLASTN	925	1e-68	85
215	1	700096890H1	SATMON008	g1778820	BLASTN	926	1e-68	90
216	1	700154404H1	SATMON007	g450548	BLASTN	928	1e-68	89
217	1	700468337H1	SATMON025	g450548	BLASTN	445	1e-67	88
218	1	700158827H1	SATMON012	g1778820	BLASTN	522	1e-67	87
219	1	700241744H1	SATMON010	g1778820	BLASTN	575	1e-67	85
220	1	700624633H1	SATMON034	g960356	BLASTN	644	1e-67	85
221	1	700154564H1	SATMON007	g1778820	BLASTN	735	1e-67	85
222	1	700172424H1	SATMON013	g450548	BLASTN	912	1e-67	90
223	1	700096287H1	SATMON008	g1778820	BLASTN	914	1e-67	86
224	1	700207638H1	SATMON016	g1778820	BLASTN	914	1e-67	86
225	1	700093735H1	SATMON008	g960356	BLASTN	920	1e-67	89
226	1	700159494H1	SATMON012	g1778820	BLASTN	899	1e-66	89
227	1	700236013H1	SATMON010	g1778820	BLASTN	900	1e-66	83
228	1	700220267H1	SATMON011	g1778820	BLASTN	439	1e-65	84
229	1	700477578H1	SATMON025	g450548	BLASTN	576	1e-65	90
230	1	700047743H1	SATMON003	g1778820	BLASTN	601	1e-65	82
231	1	700343041H1	SATMON021	g1778820	BLASTN	633	1e-65	86
232	1	700159886H1	SATMON012	g450548	BLASTN	803	1e-65	85
233	1	700570242H1	SATMON030	g450548	BLASTN	834	1e-65	85
234	1	700021930H1	SATMON001	g1778820	BLASTN	888	1e-65	87
235	1	700454959H1	SATMON029	g450548	BLASTN	890	1e-65	89
236	1	700171336H1	SATMON013	g1778820	BLASTN	890	1e-65	89
237	1	700105361H1	SATMON010	g1778820	BLASTN	806	1e-64	87
238	1	LIB3068-031-Q1-K1-B1	LIB3068	g450548	BLASTN	853	1e-64	89
239	1	700236324H1	SATMON010	g450548	BLASTN	875	1e-64	89
240	1	700150620H1	SATMON007	g450548	BLASTN	880	1e-64	87
241	1	700220648H1	SATMON011	g450548	BLASTN	881	1e-64	90
242	1	700150733H1	SATMON007	g450548	BLASTN	881	1e-64	85
243	1	700157367H1	SATMON012	g1778820	BLASTN	882	1e-64	85
244	1	700259676H1	SATMON017	g1778820	BLASTN	885	1e-64	88
245	1	700616490H1	SATMON033	g450548	BLASTN	886	1e-64	82
246	1	700102511H1	SATMON010	g450548	BLASTN	695	1e-63	89
247	1	700202805H1	SATMON003	g1778820	BLASTN	817	1e-63	92
248	1	700105685H1	SATMON010	g1778820	BLASTN	866	1e-63	84
249	1	700106113H1	SATMON010	g450548	BLASTN	873	1e-63	91
250	1	700444778H1	SATMON027	g1778820	BLASTN	343	1e-62	84
251	1	700103584H1	SATMON010	g450548	BLASTN	521	1e-62	86
252	1	LIB189-008-Q1-E1-D9	LIB189	g1778820	BLASTN	850	1e-62	91
253	1	700155684H1	SATMON007	g1778820	BLASTN	852	1e-62	85
254	1	700261639H1	SATMON017	g1778820	BLASTN	853	1e-62	89
255	1	700158367H1	SATMON012	g450548	BLASTN	856	1e-62	81
256	1	700153242H1	SATMON007	g1778820	BLASTN	859	1e-62	90
257	1	700210738H1	SATMON016	g1778820	BLASTN	859	1e-62	90
258	1	700203008H1	SATMON003	g450548	BLASTN	698	1e-61	86
259	1	700206081H1	SATMON003	g1778820	BLASTN	840	1e-61	92
260	1	700028643H1	SATMON003	g1778820	BLASTN	846	1e-61	85
261	1	700223914H1	SATMON011	g1778820	BLASTN	849	1e-61	84
262	1	700571455H1	SATMON030	g1778820	BLASTN	378	1e-60	88
263	1	700075374H1	SATMON007	g1778820	BLASTN	830	1e-60	85
264	1	700150452H1	SATMON007	g450548	BLASTN	831	1e-60	89

265	1	700261318H1	SATMON017	g1778820	BLASTN	831	1e-60	83
266	1	700616390H1	SATMON033	g450548	BLASTN	835	1e-60	92
267	1	700208718H1	SATMON016	g450548	BLASTN	561	1e-59	89
268	1	700448948H1	SATMON028	g1778820	BLASTN	653	1e-59	83
269	1	LIB3060-035-Q1-K1-H3	LIB3060	g450548	BLASTN	734	1e-59	85
270	1	700049753H1	SATMON003	g450548	BLASTN	769	1e-59	90
271	1	700154489H1	SATMON007	g1778820	BLASTN	814	1e-59	83
272	1	700170783H1	SATMON013	g1778820	BLASTN	816	1e-59	87
273	1	700237571H1	SATMON010	g450548	BLASTN	817	1e-59	89
274	1	700154872H1	SATMON007	g1778820	BLASTN	822	1e-59	85
275	1	700025620H1	SATMON004	g1778820	BLASTN	686	1e-58	84
276	1	700158255H1	SATMON012	g450548	BLASTN	803	1e-58	86
277	1	700159282H1	SATMON012	g450548	BLASTN	805	1e-58	83
278	1	700222171H1	SATMON011	g1778820	BLASTN	807	1e-58	89
279	1	700205003H1	SATMON003	g1778820	BLASTN	441	1e-57	86
280	1	700049209H1	SATMON003	g1778820	BLASTN	443	1e-57	90
281	1	700016430H1	SATMON001	g1778820	BLASTN	492	1e-57	86
282	1	700212936H1	SATMON016	g450548	BLASTN	564	1e-57	86
283	1	700156939H1	SATMON012	g1778820	BLASTN	801	1e-57	84
284	1	700222183H1	SATMON011	g1778820	BLASTN	561	1e-56	83
285	1	700203453H1	SATMON003	g1778820	BLASTN	784	1e-56	90
286	1	700167708H1	SATMON013	g1778820	BLASTN	560	1e-55	83
287	1	700214356H1	SATMON016	g1778820	BLASTN	693	1e-55	87
288	1	700475377H1	SATMON025	g450548	BLASTN	705	1e-55	90
289	1	700029625H1	SATMON003	g450548	BLASTN	712	1e-55	88
290	1	700202091H1	SATMON003	g1778820	BLASTN	774	1e-55	90
291	1	700264594H1	SATMON017	g450548	BLASTN	775	1e-55	89
292	1	700074969H1	SATMON007	g450548	BLASTN	777	1e-55	87
293	1	LIB3068-005-Q1-K1-A9	LIB3068	g1778820	BLASTN	389	1e-54	80
294	1	700207147H1	SATMON017	g1778820	BLASTN	537	1e-54	88
295	1	LIB189-004-Q1-E1-F11	LIB189	g960356	BLASTN	757	1e-54	87
296	1	700171222H1	SATMON013	g1778820	BLASTN	757	1e-54	93
297	1	700239903H1	SATMON010	g1778820	BLASTN	760	1e-54	84
298	1	700048208H1	SATMON003	g1778820	BLASTN	762	1e-54	90
299	1	700264085H1	SATMON017	g450548	BLASTN	764	1e-54	87
300	1	700155761H1	SATMON007	g1778820	BLASTN	579	1e-53	85
301	1	700216516H1	SATMON016	g450548	BLASTN	380	1e-52	89
302	1	LIB3068-044-Q1-K1-F12	LIB3068	g450548	BLASTN	492	1e-52	73
303	1	700344420H1	SATMON021	g450548	BLASTN	534	1e-52	79
304	1	700219767H1	SATMON011	g450548	BLASTN	732	1e-52	88
305	1	700153757H1	SATMON007	g1778820	BLASTN	737	1e-52	89
306	1	700165841H1	SATMON013	g450548	BLASTN	738	1e-52	90
307	1	700381966H1	SATMON023	g1778820	BLASTN	740	1e-52	86
308	1	700170427H1	SATMON013	g450548	BLASTN	687	1e-51	89
309	1	700094344H1	SATMON008	g1778820	BLASTN	725	1e-51	93
310	1	700052248H1	SATMON003	g1778820	BLASTN	726	1e-51	84
311	1	700050628H1	SATMON003	g450548	BLASTN	726	1e-51	89
312	1	700223031H1	SATMON011	g1778820	BLASTN	729	1e-51	89
313	1	700475686H1	SATMON025	g450548	BLASTN	660	1e-50	89
314	1	700170325H1	SATMON013	g2305013	BLASTN	709	1e-50	82
315	1	700221107H1	SATMON011	g450548	BLASTN	698	1e-49	89
316	1	700612589H1	SATMON033	g960356	BLASTN	703	1e-49	89
317	1	700153770H1	SATMON007	g1778820	BLASTN	705	1e-49	88
318	1	LIB3078-006-Q1-K1-E7	LIB3078	g1778820	BLASTN	705	1e-49	79

319	1	700239724H1	SATMON010	g1778820	BLASTN	507	1e-48	78
320	1	700356777H1	SATMON024	g1778820	BLASTN	683	1e-48	86
321	1	700241568H1	SATMON010	g960356	BLASTN	691	1e-48	88
322	1	700051294H1	SATMON003	g1778820	BLASTN	462	1e-47	88
323	1	700343661H1	SATMON021	g450548	BLASTN	507	1e-47	81
324	1	700029419H1	SATMON003	g1778820	BLASTN	547	1e-47	87
325	1	700165936H1	SATMON013	g1778820	BLASTN	671	1e-47	83
326	1	700026152H1	SATMON003	g960356	BLASTN	673	1e-47	89
327	1	700075671H1	SATMON007	g450548	BLASTN	429	1e-46	89
328	1	700156674H1	SATMON012	g1778820	BLASTN	659	1e-46	91
329	1	700094981H1	SATMON008	g1778820	BLASTN	662	1e-46	85
330	1	700092879H1	SATMON008	g1778820	BLASTN	668	1e-46	87
331	1	700240685H1	SATMON010	g1778820	BLASTN	484	1e-45	84
332	1	700150286H1	SATMON007	g1778820	BLASTN	647	1e-45	82
333	1	700104990H1	SATMON010	g450548	BLASTN	652	1e-45	89
334	1	700203829H1	SATMON003	g450548	BLASTN	652	1e-45	87
335	1	700153718H1	SATMON007	g167961	BLASTN	656	1e-45	91
336	1	700050841H1	SATMON003	g450548	BLASTN	571	1e-44	86
337	1	700268037H1	SATMON017	g2305013	BLASTN	636	1e-44	86
338	1	700153630H1	SATMON007	g1778820	BLASTN	637	1e-44	82
339	1	700475317H1	SATMON025	g450548	BLASTN	530	1e-43	87
340	1	701163127H1	SATMONN04	g960356	BLASTN	626	1e-43	88
341	1	700203688H1	SATMON003	g960356	BLASTN	627	1e-43	89
342	1	700049893H1	SATMON003	g1778820	BLASTN	506	1e-42	90
343	1	700449155H1	SATMON028	g1778820	BLASTN	443	1e-41	85
344	1	701183780H1	SATMONN06	g450548	BLASTN	558	1e-41	86
345	1	700242162H1	SATMON010	g1778820	BLASTN	600	1e-41	90
346	1	700466994H1	SATMON025	g450548	BLASTN	604	1e-41	91
347	1	700259823H1	SATMON017	g450548	BLASTN	607	1e-41	83
348	1	700346242H1	SATMON021	g450548	BLASTN	608	1e-41	86
349	1	700156395H1	SATMON007	g1778820	BLASTN	609	1e-41	90
350	1	700236835H1	SATMON010	g1778820	BLASTN	335	1e-40	81
351	1	700172385H1	SATMON013	g1778820	BLASTN	397	1e-40	87
352	1	700210466H1	SATMON016	g1778820	BLASTN	586	1e-40	81
353	1	700257303H1	SATMON017	g1778820	BLASTN	589	1e-40	81
354	1	LIB3067-027-Q1-K1-G1	LIB3067	g1778820	BLASTN	623	1e-40	87
355	1	LIB3066-025-Q1-K1-D1	LIB3066	g450548	BLASTN	524	1e-39	85
356	1	700106853H1	SATMON010	g450548	BLASTN	580	1e-39	86
357	1	700160540H1	SATMON012	g960356	BLASTN	581	1e-39	88
358	1	700157780H1	SATMON012	g960356	BLASTN	581	1e-39	88
359	1	700149801H1	SATMON007	g450548	BLASTN	565	1e-38	86
360	1	700353243H1	SATMON024	g450548	BLASTN	570	1e-38	86
361	1	700166171H1	SATMON013	g1778820	BLASTN	254	1e-37	79
362	1	700142509H1	SATMON012	g960356	BLASTN	556	1e-37	88
363	1	700242131H1	SATMON010	g450548	BLASTN	560	1e-37	85
364	1	700455643H1	SATMON029	g450548	BLASTN	539	1e-36	86
365	1	700150248H1	SATMON007	g1778820	BLASTN	547	1e-36	87
366	1	700208549H1	SATMON016	g450548	BLASTN	559	1e-36	88
367	1	700027193H1	SATMON003	g450548	BLASTN	529	1e-35	86
368	1	700221390H1	SATMON011	g450548	BLASTN	530	1e-35	85
369	1	700455647H1	SATMON029	g450548	BLASTN	531	1e-35	85
370	1	700260103H1	SATMON017	g1778820	BLASTN	531	1e-35	80
371	1	700167344H1	SATMON013	g1778820	BLASTN	536	1e-35	88
372	1	700089913H1	SATMON011	g960356	BLASTN	546	1e-35	88

373	1	700570573H1	SATMON030	g450548	BLASTN	300	1e-34	89
374	1	700169889H1	SATMON013	g1778820	BLASTN	519	1e-34	89
375	1	700262857H1	SATMON017	g1778820	BLASTN	521	1e-34	87
376	1	700142644H2	SATMON013	g1778820	BLASTN	522	1e-34	87
377	1	700224417H1	SATMON011	g450548	BLASTN	522	1e-34	91
378	1	700073882H1	SATMON007	g450548	BLASTN	531	1e-34	87
379	1	700085803H1	SATMON011	g450548	BLASTN	338	1e-33	89
380	1	700162323H1	SATMON012	g1778820	BLASTN	513	1e-33	86
381	1	700468306H1	SATMON025	g450548	BLASTN	519	1e-33	89
382	1	700443224H2	SATMON026	g450548	BLASTN	275	1e-32	83
383	1	LIB3066-054-Q1-K1-E3	LIB3066	g450548	BLASTN	495	1e-32	87
384	1	700211827H1	SATMON016	g1778820	BLASTN	489	1e-31	80
385	1	700048741H1	SATMON003	g450548	BLASTN	500	1e-31	88
386	1	700613620H1	SATMON033	g1778820	BLASTN	385	1e-30	88
387	1	700029203H1	SATMON003	g1778820	BLASTN	461	1e-29	84
388	1	700378431H1	SATMON020	g450548	BLASTN	466	1e-29	89
389	1	700455641H1	SATMON029	g450548	BLASTN	472	1e-29	85
390	1	700447867H1	SATMON027	g960356	BLASTN	473	1e-29	88
391	1	700447511H1	SATMON027	g450548	BLASTN	474	1e-29	88
392	1	700025851H1	SATMON003	g450548	BLASTN	479	1e-29	88
393	1	LIB3066-024-Q1-K1-H4	LIB3066	g1778820	BLASTN	481	1e-29	81
394	1	LIB143-025-Q1-E1-C4	LIB143	g450549	BLASTX	64	1e-28	70
395	1	700242282H1	SATMON010	g1778820	BLASTN	280	1e-28	85
396	1	700087244H1	SATMON011	g450548	BLASTN	464	1e-28	88
397	1	700458127H1	SATMON029	g450548	BLASTN	238	1e-27	82
398	1	700025767H1	SATMON003	g1778820	BLASTN	438	1e-26	86
399	1	700235401H1	SATMON010	g450548	BLASTN	442	1e-26	88
400	1	700029457H1	SATMON003	g450548	BLASTN	446	1e-26	89
401	1	700266174H1	SATMON017	g450548	BLASTN	447	1e-26	89
402	1	700349254H1	SATMON023	g1778820	BLASTN	315	1e-25	85
403	1	LIB3068-041-Q1-K1-H6	LIB3068	g450548	BLASTN	438	1e-25	86
404	1	700092889H1	SATMON008	g1778820	BLASTN	397	1e-24	92
405	1	700049044H1	SATMON003	g1778820	BLASTN	425	1e-24	90
406	1	700202710H1	SATMON003	g960357	BLASTX	114	1e-19	97
407	1	700155117H1	SATMON007	g450548	BLASTN	314	1e-19	90
408	1	700449619H1	SATMON028	g1778820	BLASTN	341	1e-19	89
409	1	700150336H1	SATMON007	g1778820	BLASTN	343	1e-19	87
410	1	700166223H1	SATMON013	g960357	BLASTX	181	1e-18	97
411	1	700153796H1	SATMON007	g2305013	BLASTN	300	1e-16	82
412	1	700053266H1	SATMON008	g450548	BLASTN	292	1e-15	83
413	1	700405227H1	SATMON028	g450548	BLASTN	313	1e-15	90
414	1	700397410H1	SATMONN01	g17262	BLASTX	147	1e-14	96
415	1	700211524H1	SATMON016	g1033190	BLASTX	153	1e-14	100
416	1	700281415H2	SATMON019	g2315140	BLASTX	139	1e-13	89
417	1	700429873H1	SATMONN01	g16961	BLASTX	133	1e-12	94
418	1	700239660H1	SATMON010	g450548	BLASTN	245	1e-12	86
419	1	700213596H1	SATMON016	g450548	BLASTN	258	1e-12	95
420	1	700152367H1	SATMON007	g450548	BLASTN	273	1e-12	88
421	1	700452014H1	SATMON028	g17262	BLASTX	76	1e-11	72
422	1	700357106H1	SATMON024	g1724104	BLASTX	132	1e-11	100
423	1	700468611H1	SATMON025	g450549	BLASTX	93	1e-10	93
424	1	700213526H1	SATMON016	g450549	BLASTX	127	1e-10	96
425	1	700267065H1	SATMON017	g450549	BLASTX	130	1e-10	96
426	1	700152044H1	SATMON007	g450549	BLASTX	88	1e-8	93

427	1	700159090H1	SATMON012	g169665	BLASTX	113	1e-8	81
428	1	700266734H1	SATMON017	g450549	BLASTX	119	1e-8	96
429	1	700405367H1	SATMON029	g1778820	BLASTN	231	1e-8	65
1635	-700555532	700555532H1	SOYMON001	g429103	BLASTN	920	1e-67	84
1636	-700649594	700649594H1	SOYMON003	g609559	BLASTX	186	1e-23	85
1637	-700750590	700750590H1	SOYMON014	g609224	BLASTN	363	1e-40	79
1638	-700755802	700755802H1	SOYMON014	g609224	BLASTN	479	1e-43	74
1639	-700869211	700869211H1	SOYMON016	g169665	BLASTX	146	1e-13	92
1640	-700891960	700891960H1	SOYMON024	g726031	BLASTN	589	1e-56	82
1641	-700900377	700900377H1	SOYMON027	g1655577	BLASTN	235	1e-8	78
1642	-700902427	700902427H1	SOYMON027	g1655576	BLASTX	151	1e-13	76
1643	-700941686	700941686H1	SOYMON024	g497899	BLASTN	442	1e-26	89
1644	-700952418	700952418H1	SOYMON022	g726030	BLASTX	146	1e-17	83
1645	-700979651	700979651H2	SOYMON009	g1655579	BLASTN	1006	1e-75	84
1646	-700982809	700982809H1	SOYMON009	g1655579	BLASTN	945	1e-69	82
1647	-700982867	700982867H1	SOYMON009	g1127582	BLASTN	868	1e-63	78
1648	-701056884	701056884H1	SOYMON032	g609556	BLASTN	726	1e-51	84
1649	-701117318	701117318H1	SOYMON037	g609224	BLASTN	451	1e-37	80
1650	-701118224	701118224H1	SOYMON037	g2305013	BLASTN	285	1e-19	72
1651	-701121264	701121264H1	SOYMON037	g166873	BLASTN	238	1e-8	82
1652	-701122908	701122908H1	SOYMON037	g16508	BLASTN	444	1e-26	83
1653	-701128589	701128589H1	SOYMON037	g16508	BLASTN	539	1e-36	90
1654	-GM12798	LIB3049-039-Q1-E1-F2	LIB3049	g16508	BLASTN	578	1e-37	73
1655	-GM14331	LIB3049-055-Q1-E1-F5	LIB3049	g167961	BLASTN	497	1e-30	62
1656	-GM30881	LIB3050-005-Q1-K1-G1	LIB3050	g1655577	BLASTN	543	1e-34	82
1657	-GM30911	LIB3050-005-Q1-K1-B8	LIB3050	g1655577	BLASTN	387	1e-26	71
1658	-GM33921	LIB3051-028-Q1-K1-A9	LIB3051	g16508	BLASTN	338	1e-32	73
1659	12644	701131794H1	SOYMON038	g429107	BLASTN	877	1e-64	84
1660	12644	701142515H1	SOYMON038	g429107	BLASTN	871	1e-63	84
1661	12644	700888494H1	SOYMON024	g429107	BLASTN	662	1e-46	83
1662	16	LIB3030-009-Q1-B1-C1	LIB3030	g429105	BLASTN	1439	1e-119	84
1663	16	LIB3051-106-Q1-K1-B5	LIB3051	g609224	BLASTN	1516	1e-117	86
1664	16	LIB3050-023-Q1-K1-A12	LIB3050	g1724103	BLASTN	1388	1e-106	83
1665	16	LIB3028-003-Q1-B1-G11	LIB3028	g609224	BLASTN	1313	1e-100	87
1666	16	LIB3054-010-Q1-N1-E2	LIB3054	g609224	BLASTN	920	1e-95	87
1667	16	700651294H1	SOYMON003	g609224	BLASTN	672	1e-94	86
1668	16	LIB3027-007-Q1-B1-G3	LIB3027	g16508	BLASTN	891	1e-93	83
1669	16	LIB3053-013-Q1-N1-H9	LIB3053	g609224	BLASTN	996	1e-93	87
1670	16	LIB3051-061-Q1-K1-C7	LIB3051	g16508	BLASTN	1195	1e-93	86
1671	16	LIB3065-005-Q1-N1-A6	LIB3065	g609224	BLASTN	707	1e-91	78
1672	16	LIB3050-008-Q1-E1-E3	LIB3050	g609224	BLASTN	1102	1e-87	87
1673	16	LIB3051-011-Q1-E1-B2	LIB3051	g16508	BLASTN	1153	1e-87	82
1674	16	LIB3030-010-Q1-B1-F8	LIB3030	g609224	BLASTN	988	1e-86	83
1675	16	700652104H1	SOYMON003	g1655577	BLASTN	990	1e-86	83
1676	16	701205279H1	SOYMON035	g609556	BLASTN	1125	1e-85	86
1677	16	700662183H1	SOYMON005	g609224	BLASTN	678	1e-83	84
1678	16	LIB3051-039-Q1-K1-F5	LIB3051	g169664	BLASTN	1104	1e-83	86
1679	16	700557616H1	SOYMON001	g609556	BLASTN	1111	1e-83	86
1680	16	LIB3030-002-Q1-B1-C6	LIB3030	g16508	BLASTN	1113	1e-83	83
1681	16	700865235H1	SOYMON016	g1724103	BLASTN	1090	1e-82	84
1682	16	700563340H1	SOYMON002	g609224	BLASTN	911	1e-80	86
1683	16	LIB3040-044-Q1-E1-D7	LIB3040	g166873	BLASTN	1052	1e-80	82
1684	16	LIB3040-060-Q1-E1-D9	LIB3040	g609224	BLASTN	1071	1e-80	84
1685	16	LIB3049-001-Q1-E1-F12	LIB3049	g16508	BLASTN	1074	1e-80	84

1686	16	LIB3028-006-Q1-B1-F1	LIB3028	g16508	BLASTN	857	1e-79	83
1687	16	LIB3049-033-Q1-E1-G5	LIB3049	g2315139	BLASTN	933	1e-79	81
1688	16	700978240H1	SOYMON009	g609224	BLASTN	984	1e-79	88
1689	16	700980802H1	SOYMON009	g862999	BLASTN	1060	1e-79	85
1690	16	700755908H1	SOYMON014	g1724103	BLASTN	1062	1e-79	88
1691	16	700562226H1	SOYMON002	g1724103	BLASTN	1065	1e-79	84
1692	16	701119101H1	SOYMON037	g609224	BLASTN	1048	1e-78	86
1693	16	700646291H1	SOYMON012	g1655577	BLASTN	1049	1e-78	86
1694	16	LIB3050-022-Q1-K1-B9	LIB3050	g16508	BLASTN	1049	1e-78	83
1695	16	LIB3028-030-Q1-B1-F8	LIB3028	g16508	BLASTN	1053	1e-78	83
1696	16	701143128H1	SOYMON038	g609556	BLASTN	737	1e-77	86
1697	16	LIB3040-041-Q1-E1-D10	LIB3040	g166873	BLASTN	861	1e-77	81
1698	16	700756363H1	SOYMON014	g609556	BLASTN	1031	1e-77	88
1699	16	700729963H1	SOYMON009	g1724103	BLASTN	1036	1e-77	88
1700	16	701063046H1	SOYMON033	g609224	BLASTN	1037	1e-77	86
1701	16	LIB3030-005-Q1-B1-G2	LIB3030	g609224	BLASTN	1038	1e-77	84
1702	16	700564331H1	SOYMON002	g609556	BLASTN	1039	1e-77	84
1703	16	700562958H1	SOYMON002	g1724103	BLASTN	1040	1e-77	88
1704	16	LIB3039-014-Q1-E1-F7	LIB3039	g16508	BLASTN	569	1e-76	84
1705	16	700753632H1	SOYMON014	g497899	BLASTN	668	1e-76	83
1706	16	LIB3049-048-Q1-E1-F2	LIB3049	g16508	BLASTN	722	1e-76	83
1707	16	700648911H1	SOYMON003	g1655577	BLASTN	983	1e-76	83
1708	16	LIB3040-027-Q1-E1-H3	LIB3040	g166873	BLASTN	1003	1e-76	81
1709	16	700664905H1	SOYMON005	g609556	BLASTN	1020	1e-76	86
1710	16	701133404H1	SOYMON038	g1724103	BLASTN	1023	1e-76	85
1711	16	701208780H1	SOYMON035	g1655577	BLASTN	1025	1e-76	85
1712	16	700902279H1	SOYMON027	g169664	BLASTN	1026	1e-76	88
1713	16	LIB3040-048-Q1-E1-G10	LIB3040	g16508	BLASTN	1030	1e-76	83
1714	16	LIB3040-028-Q1-E1-A4	LIB3040	g16508	BLASTN	979	1e-75	84
1715	16	700807586H1	SOYMON016	g609224	BLASTN	1006	1e-75	86
1716	16	700755486H1	SOYMON014	g609224	BLASTN	1006	1e-75	87
1717	16	700724920H1	SOYMON009	g609224	BLASTN	1009	1e-75	87
1718	16	701007494H2	SOYMON019	g1724103	BLASTN	1013	1e-75	85
1719	16	700568310H1	SOYMON002	g497899	BLASTN	834	1e-74	83
1720	16	701208819H1	SOYMON035	g609224	BLASTN	857	1e-74	87
1721	16	700985084H1	SOYMON009	g609224	BLASTN	872	1e-74	85
1722	16	701013074H1	SOYMON019	g726031	BLASTN	894	1e-74	88
1723	16	700650843H1	SOYMON003	g609224	BLASTN	924	1e-74	86
1724	16	700646037H1	SOYMON011	g609556	BLASTN	930	1e-74	84
1725	16	LIB3040-039-Q1-E1-H9	LIB3040	g166873	BLASTN	973	1e-74	80
1726	16	700847231H1	SOYMON021	g726031	BLASTN	999	1e-74	85
1727	16	700792293H1	SOYMON011	g609224	BLASTN	999	1e-74	86
1728	16	701109644H1	SOYMON036	g609556	BLASTN	1004	1e-74	86
1729	16	701122929H1	SOYMON037	g1724103	BLASTN	459	1e-73	87
1730	16	700661491H1	SOYMON005	g16508	BLASTN	540	1e-73	90
1731	16	701120070H1	SOYMON037	g609224	BLASTN	786	1e-73	87
1732	16	700898895H1	SOYMON027	g609556	BLASTN	820	1e-73	88
1733	16	701096959H1	SOYMON028	g429105	BLASTN	839	1e-73	82
1734	16	700848356H1	SOYMON021	g497899	BLASTN	985	1e-73	85
1735	16	700864876H1	SOYMON016	g1724103	BLASTN	985	1e-73	88
1736	16	700868456H1	SOYMON016	g609224	BLASTN	987	1e-73	86
1737	16	701210488H1	SOYMON035	g1655577	BLASTN	988	1e-73	83
1738	16	700747764H1	SOYMON013	g609224	BLASTN	822	1e-72	87
1739	16	701063311H1	SOYMON033	g1655577	BLASTN	971	1e-72	86

1740	16	700868625H1	SOYMON016	g1724103	BLASTN	973	1e-72	85
1741	16	700992344H1	SOYMON011	g726031	BLASTN	973	1e-72	85
1742	16	701012719H1	SOYMON019	g1724103	BLASTN	973	1e-72	85
1743	16	700676769H1	SOYMON007	g609224	BLASTN	973	1e-72	87
1744	16	700946235H1	SOYMON024	g609224	BLASTN	975	1e-72	85
1745	16	700891218H1	SOYMON024	g1724103	BLASTN	975	1e-72	84
1746	16	700724907H1	SOYMON009	g609556	BLASTN	977	1e-72	84
1747	16	700833549H1	SOYMON019	g1655577	BLASTN	979	1e-72	85
1748	16	700942105H1	SOYMON024	g726031	BLASTN	980	1e-72	84
1749	16	700564290H1	SOYMON002	g726031	BLASTN	530	1e-71	84
1750	16	700891233H1	SOYMON024	g1724103	BLASTN	746	1e-71	86
1751	16	LIB3028-025-Q1-B1-B2	LIB3028	g609224	BLASTN	870	1e-71	85
1752	16	701120896H1	SOYMON037	g429105	BLASTN	959	1e-71	82
1753	16	LIB3051-027-Q1-K1-A9	LIB3051	g609224	BLASTN	962	1e-71	83
1754	16	701050696H1	SOYMON032	g609556	BLASTN	965	1e-71	86
1755	16	700653619H1	SOYMON003	g609224	BLASTN	966	1e-71	86
1756	16	701047994H1	SOYMON032	g1724103	BLASTN	967	1e-71	85
1757	16	701123056H1	SOYMON037	g1724103	BLASTN	968	1e-71	85
1758	16	700983479H1	SOYMON009	g726031	BLASTN	653	1e-70	85
1759	16	701063733H1	SOYMON034	g1655577	BLASTN	850	1e-70	85
1760	16	700738366H1	SOYMON012	g726031	BLASTN	946	1e-70	85
1761	16	700866319H1	SOYMON016	g1655575	BLASTN	947	1e-70	85
1762	16	700789013H2	SOYMON011	g1655577	BLASTN	948	1e-70	85
1763	16	700896080H1	SOYMON027	g609224	BLASTN	949	1e-70	87
1764	16	LIB3039-021-Q1-E1-F12	LIB3039	g16508	BLASTN	950	1e-70	83
1765	16	700898284H1	SOYMON027	g429105	BLASTN	950	1e-70	86
1766	16	700901743H1	SOYMON027	g609224	BLASTN	950	1e-70	86
1767	16	700831048H1	SOYMON019	g429105	BLASTN	951	1e-70	85
1768	16	701038107H1	SOYMON029	g726031	BLASTN	952	1e-70	87
1769	16	700559703H1	SOYMON001	g726031	BLASTN	954	1e-70	88
1770	16	700848817H1	SOYMON021	g609224	BLASTN	954	1e-70	84
1771	16	700896032H1	SOYMON027	g609224	BLASTN	956	1e-70	87
1772	16	700944764H1	SOYMON024	g16508	BLASTN	567	1e-69	85
1773	16	701011015H1	SOYMON019	g1724103	BLASTN	647	1e-69	81
1774	16	701125662H1	SOYMON037	g609224	BLASTN	751	1e-69	87
1775	16	701046895H1	SOYMON032	g16508	BLASTN	795	1e-69	83
1776	16	701012632H1	SOYMON019	g1655577	BLASTN	800	1e-69	85
1777	16	701005244H1	SOYMON019	g1724103	BLASTN	804	1e-69	82
1778	16	LIB3050-023-Q1-K1-H10	LIB3050	g609224	BLASTN	899	1e-69	83
1779	16	700892558H1	SOYMON024	g1655575	BLASTN	936	1e-69	85
1780	16	700988779H1	SOYMON011	g16508	BLASTN	937	1e-69	82
1781	16	701203923H2	SOYMON035	g429105	BLASTN	938	1e-69	86
1782	16	701010231H2	SOYMON019	g1724103	BLASTN	938	1e-69	83
1783	16	701041790H1	SOYMON029	g450548	BLASTN	939	1e-69	84
1784	16	700967887H1	SOYMON033	g1724103	BLASTN	940	1e-69	87
1785	16	701123361H1	SOYMON037	g16508	BLASTN	941	1e-69	84
1786	16	701123154H1	SOYMON037	g1724103	BLASTN	942	1e-69	86
1787	16	701045767H1	SOYMON032	g726031	BLASTN	942	1e-69	85
1788	16	700983288H1	SOYMON009	g609224	BLASTN	945	1e-69	83
1789	16	700653053H1	SOYMON003	g609224	BLASTN	945	1e-69	86
1790	16	701044226H1	SOYMON032	g1655577	BLASTN	486	1e-68	83
1791	16	700969927H1	SOYMON005	g497899	BLASTN	750	1e-68	85
1792	16	700547956H1	SOYMON001	g609224	BLASTN	779	1e-68	87
1793	16	LIB3049-048-Q1-E1-E4	LIB3049	g609224	BLASTN	827	1e-68	83

1794	16	701140780H1	SOYMON038	g1655577	BLASTN	922	1e-68	85
1795	16	700849166H1	SOYMON021	g1724103	BLASTN	923	1e-68	84
1796	16	700891945H1	SOYMON024	g609556	BLASTN	925	1e-68	87
1797	16	700658051H1	SOYMON004	g429105	BLASTN	925	1e-68	83
1798	16	700942415H1	SOYMON024	g1655575	BLASTN	928	1e-68	84
1799	16	701041890H1	SOYMON029	g1724103	BLASTN	930	1e-68	82
1800	16	LIB3028-006-Q1-B1-H12	LIB3028	g16508	BLASTN	931	1e-68	81
1801	16	700967039H1	SOYMON029	g429105	BLASTN	931	1e-68	85
1802	16	700836178H1	SOYMON019	g1724103	BLASTN	933	1e-68	85
1803	16	700897558H1	SOYMON027	g16508	BLASTN	737	1e-67	85
1804	16	LIB3050-004-Q1-E1-A2	LIB3050	g609224	BLASTN	796	1e-67	80
1805	16	700730236H1	SOYMON009	g726031	BLASTN	816	1e-67	84
1806	16	700833936H1	SOYMON019	g1724103	BLASTN	915	1e-67	84
1807	16	700897552H1	SOYMON027	g1655577	BLASTN	916	1e-67	84
1808	16	700945440H1	SOYMON024	g609556	BLASTN	917	1e-67	84
1809	16	700961368H1	SOYMON022	g169664	BLASTN	918	1e-67	88
1810	16	700789702H1	SOYMON011	g609224	BLASTN	921	1e-67	86
1811	16	700786541H1	SOYMON011	g429105	BLASTN	513	1e-66	84
1812	16	700987282H1	SOYMON009	g16508	BLASTN	523	1e-66	87
1813	16	700661002H1	SOYMON005	g2305013	BLASTN	836	1e-66	81
1814	16	701148312H1	SOYMON031	g862999	BLASTN	899	1e-66	84
1815	16	700738822H1	SOYMON012	g1655577	BLASTN	899	1e-66	85
1816	16	700940996H1	SOYMON024	g609556	BLASTN	901	1e-66	86
1817	16	700749195H1	SOYMON013	g609556	BLASTN	903	1e-66	81
1818	16	700893941H1	SOYMON024	g429105	BLASTN	903	1e-66	88
1819	16	700892888H1	SOYMON024	g609556	BLASTN	906	1e-66	86
1820	16	700901481H1	SOYMON027	g169664	BLASTN	638	1e-65	86
1821	16	700945269H1	SOYMON024	g169664	BLASTN	688	1e-65	86
1822	16	700746876H1	SOYMON013	g609556	BLASTN	740	1e-65	85
1823	16	700755043H1	SOYMON014	g609556	BLASTN	760	1e-65	87
1824	16	701097166H1	SOYMON028	g1655577	BLASTN	781	1e-65	82
1825	16	701129484H1	SOYMON037	g16508	BLASTN	898	1e-65	82
1826	16	700651014H1	SOYMON003	g167961	BLASTN	464	1e-64	81
1827	16	701134363H1	SOYMON038	g726031	BLASTN	687	1e-64	84
1828	16	701124677H1	SOYMON037	g726031	BLASTN	876	1e-64	85
1829	16	701000359H1	SOYMON018	g1724103	BLASTN	877	1e-64	87
1830	16	701139375H1	SOYMON038	g1724103	BLASTN	883	1e-64	84
1831	16	700832784H1	SOYMON019	g609224	BLASTN	883	1e-64	89
1832	16	700980227H1	SOYMON009	g1724103	BLASTN	884	1e-64	84
1833	16	700943177H1	SOYMON024	g1655577	BLASTN	885	1e-64	85
1834	16	700844446H1	SOYMON021	g2315139	BLASTN	355	1e-63	84
1835	16	700836453H1	SOYMON020	g862999	BLASTN	494	1e-63	85
1836	16	700983012H1	SOYMON009	g16508	BLASTN	756	1e-63	83
1837	16	700795805H1	SOYMON017	g1655577	BLASTN	833	1e-63	85
1838	16	701213663H1	SOYMON035	g1724103	BLASTN	862	1e-63	84
1839	16	700868366H1	SOYMON016	g167961	BLASTN	864	1e-63	83
1840	16	701134377H1	SOYMON038	g2305013	BLASTN	865	1e-63	79
1841	16	LIB3049-007-Q1-E1-C9	LIB3049	g16508	BLASTN	867	1e-63	79
1842	16	700944577H1	SOYMON024	g862999	BLASTN	869	1e-63	85
1843	16	700992289H1	SOYMON011	g16508	BLASTN	869	1e-63	84
1844	16	700891612H1	SOYMON024	g609556	BLASTN	872	1e-63	86
1845	16	700738277H1	SOYMON012	g609224	BLASTN	463	1e-62	85
1846	16	700895571H1	SOYMON027	g609224	BLASTN	562	1e-62	88
1847	16	LIB3049-008-Q1-E1-B3	LIB3049	g16508	BLASTN	853	1e-62	77



1848	16	LIB3027-010-Q1-B1-E12	LIB3027	g609224	BLASTN	854	1e-62	82
1849	16	701129389H1	SOYMON037	g16508	BLASTN	854	1e-62	83
1850	16	701045542H1	SOYMON032	g429105	BLASTN	855	1e-62	86
1851	16	700969901H1	SOYMON005	g726031	BLASTN	856	1e-62	84
1852	16	700984664H1	SOYMON009	g1724103	BLASTN	856	1e-62	81
1853	16	700561352H1	SOYMON002	g609224	BLASTN	858	1e-62	85
1854	16	701138828H1	SOYMON038	g16508	BLASTN	858	1e-62	84
1855	16	700845962H1	SOYMON021	g1655577	BLASTN	861	1e-62	87
1856	16	700896147H1	SOYMON027	g16508	BLASTN	336	1e-61	82
1857	16	701137929H1	SOYMON038	g1655577	BLASTN	502	1e-61	84
1858	16	LIB3040-032-Q1-E1-B8	LIB3040	g16508	BLASTN	521	1e-61	82
1859	16	701009537H1	SOYMON019	g726031	BLASTN	545	1e-61	86
1860	16	LIB3049-031-Q1-E1-E4	LIB3049	g609224	BLASTN	677	1e-61	82
1861	16	LIB3049-031-Q1-E1-C7	LIB3049	g167961	BLASTN	696	1e-61	82
1862	16	700891843H1	SOYMON024	g1655577	BLASTN	737	1e-61	84
1863	16	700561231H1	SOYMON002	g16508	BLASTN	839	1e-61	82
1864	16	700548238H1	SOYMON002	g429105	BLASTN	843	1e-61	85
1865	16	700901018H1	SOYMON027	g2305013	BLASTN	845	1e-61	82
1866	16	701134413H1	SOYMON038	g2305013	BLASTN	848	1e-61	82
1867	16	700653524H1	SOYMON003	g609224	BLASTN	499	1e-60	85
1868	16	LIB3049-017-Q1-E1-G8	LIB3049	g16508	BLASTN	514	1e-60	82
1869	16	701121077H1	SOYMON037	g16508	BLASTN	671	1e-60	84
1870	16	LIB3056-013-Q1-N1-A9	LIB3056	g609224	BLASTN	784	1e-60	86
1871	16	700943524H1	SOYMON024	g16508	BLASTN	831	1e-60	85
1872	16	700952889H1	SOYMON022	g429103	BLASTN	835	1e-60	84
1873	16	700730632H1	SOYMON009	g1655577	BLASTN	431	1e-59	82
1874	16	700649136H1	SOYMON003	g609224	BLASTN	486	1e-59	84
1875	16	700895591H1	SOYMON027	g609556	BLASTN	492	1e-59	87
1876	16	701009829H1	SOYMON019	g429103	BLASTN	572	1e-59	80
1877	16	LIB3039-011-Q1-E1-C11	LIB3039	g16508	BLASTN	619	1e-59	83
1878	16	LIB3040-055-Q1-E1-C4	LIB3040	g16508	BLASTN	622	1e-59	82
1879	16	701123571H1	SOYMON037	g2305013	BLASTN	682	1e-59	81
1880	16	LIB3049-004-Q1-E1-E5	LIB3049	g167961	BLASTN	771	1e-59	83
1881	16	701002239H1	SOYMON018	g609224	BLASTN	782	1e-59	86
1882	16	700971088H1	SOYMON005	g1724103	BLASTN	793	1e-59	81
1883	16	700864624H1	SOYMON016	g167961	BLASTN	817	1e-59	83
1884	16	700863534H1	SOYMON027	g16508	BLASTN	818	1e-59	86
1885	16	700749489H1	SOYMON013	g16508	BLASTN	823	1e-59	86
1886	16	700730689H1	SOYMON009	g16508	BLASTN	825	1e-59	83
1887	16	LIB3049-050-Q1-E1-A11	LIB3049	g16508	BLASTN	831	1e-59	83
1888	16	700850803H1	SOYMON023	g1655577	BLASTN	511	1e-58	83
1889	16	700992317H1	SOYMON011	g1655577	BLASTN	520	1e-58	79
1890	16	701119129H1	SOYMON037	g16508	BLASTN	553	1e-58	83
1891	16	700657623H1	SOYMON004	g1724103	BLASTN	600	1e-58	83
1892	16	LIB3049-015-Q1-E1-F8	LIB3049	g16508	BLASTN	674	1e-58	79
1893	16	701007027H1	SOYMON019	g16508	BLASTN	804	1e-58	81
1894	16	701120820H1	SOYMON037	g609224	BLASTN	807	1e-58	85
1895	16	700654317H1	SOYMON004	g1655577	BLASTN	811	1e-58	84
1896	16	700889071H1	SOYMON024	g497899	BLASTN	812	1e-58	83
1897	16	701010676H1	SOYMON019	g609224	BLASTN	813	1e-58	85
1898	16	701099590H1	SOYMON028	g1655577	BLASTN	486	1e-57	82
1899	16	700649684H1	SOYMON003	g16508	BLASTN	540	1e-57	82
1900	16	700994107H1	SOYMON011	g16508	BLASTN	567	1e-57	80
1901	16	700898629H1	SOYMON027	g16508	BLASTN	687	1e-57	82

1902	16	700902333H1	SOYMON027	g609224	BLASTN	700	1e-57	82
1903	16	LIB3039-017-Q1-E1-D9	LIB3039	g16508	BLASTN	703	1e-57	81
1904	16	LIB3040-026-Q1-E1-A4	LIB3040	g16508	BLASTN	709	1e-57	83
1905	16	701130101H1	SOYMON037	g1724103	BLASTN	791	1e-57	88
1906	16	700902482H1	SOYMON027	g169664	BLASTN	797	1e-57	81
1907	16	700730789H1	SOYMON009	g16508	BLASTN	798	1e-57	82
1908	16	700868670H1	SOYMON016	g16508	BLASTN	800	1e-57	83
1909	16	LIB3029-011-Q1-B1-D1	LIB3029	g609224	BLASTN	800	1e-57	82
1910	16	701119278H1	SOYMON037	g609224	BLASTN	801	1e-57	83
1911	16	700747731H1	SOYMON013	g16508	BLASTN	801	1e-57	86
1912	16	LIB3049-017-Q1-E1-A11	LIB3049	g16508	BLASTN	811	1e-57	80
1913	16	700890428H1	SOYMON024	g609224	BLASTN	482	1e-56	83
1914	16	700562326H1	SOYMON002	g609224	BLASTN	778	1e-56	82
1915	16	700567740H1	SOYMON002	g609224	BLASTN	781	1e-56	85
1916	16	701061514H1	SOYMON033	g1724103	BLASTN	784	1e-56	86
1917	16	701135670H1	SOYMON038	g429105	BLASTN	785	1e-56	81
1918	16	701139657H1	SOYMON038	g609224	BLASTN	786	1e-56	85
1919	16	701123371H1	SOYMON037	g16508	BLASTN	788	1e-56	86
1920	16	700838744H1	SOYMON020	g16508	BLASTN	789	1e-56	82
1921	16	701070458H1	SOYMON034	g1655577	BLASTN	457	1e-55	78
1922	16	701003437H1	SOYMON019	g429105	BLASTN	474	1e-55	83
1923	16	700835976H1	SOYMON019	g16508	BLASTN	562	1e-55	84
1924	16	700894535H1	SOYMON024	g16508	BLASTN	570	1e-55	86
1925	16	700752755H1	SOYMON014	g1655577	BLASTN	690	1e-55	83
1926	16	LIB3049-032-Q1-E1-C9	LIB3049	g16508	BLASTN	700	1e-55	83
1927	16	LIB3040-030-Q1-E1-B5	LIB3040	g16508	BLASTN	709	1e-55	84
1928	16	701136935H1	SOYMON038	g609224	BLASTN	766	1e-55	85
1929	16	700682088H1	SOYMON008	g169664	BLASTN	767	1e-55	90
1930	16	700682188H1	SOYMON008	g169664	BLASTN	767	1e-55	90
1931	16	701068649H1	SOYMON034	g16508	BLASTN	770	1e-55	83
1932	16	700726623H1	SOYMON009	g16508	BLASTN	771	1e-55	85
1933	16	700751129H1	SOYMON014	g16508	BLASTN	772	1e-55	86
1934	16	700945234H1	SOYMON024	g16508	BLASTN	773	1e-55	83
1935	16	701133507H2	SOYMON038	g609224	BLASTN	776	1e-55	85
1936	16	700902459H1	SOYMON027	g726031	BLASTN	777	1e-55	84
1937	16	700986624H1	SOYMON009	g16508	BLASTN	612	1e-54	83
1938	16	701097020H1	SOYMON028	g609224	BLASTN	615	1e-54	84
1939	16	701131409H1	SOYMON038	g16508	BLASTN	755	1e-54	91
1940	16	700750123H1	SOYMON013	g726031	BLASTN	756	1e-54	86
1941	16	701040251H1	SOYMON029	g16508	BLASTN	756	1e-54	85
1942	16	700732290H1	SOYMON010	g169664	BLASTN	758	1e-54	90
1943	16	701117458H1	SOYMON037	g609224	BLASTN	759	1e-54	82
1944	16	701118709H1	SOYMON037	g16508	BLASTN	760	1e-54	91
1945	16	701012834H1	SOYMON019	g16508	BLASTN	760	1e-54	91
1946	16	701133316H1	SOYMON038	g16508	BLASTN	760	1e-54	91
1947	16	701129646H1	SOYMON037	g16508	BLASTN	760	1e-54	91
1948	16	700732265H1	SOYMON010	g169664	BLASTN	760	1e-54	90
1949	16	701040796H1	SOYMON029	g1724103	BLASTN	760	1e-54	89
1950	16	700846350H1	SOYMON021	g16508	BLASTN	761	1e-54	86
1951	16	701137005H1	SOYMON038	g16508	BLASTN	761	1e-54	86
1952	16	701046506H1	SOYMON032	g16508	BLASTN	761	1e-54	91
1953	16	700894462H1	SOYMON024	g16508	BLASTN	761	1e-54	86
1954	16	700973847H1	SOYMON005	g16508	BLASTN	766	1e-54	87
1955	16	701128215H1	SOYMON037	g609556	BLASTN	466	1e-53	82

1956	16	700842468H1	SOYMON020	g429105	BLASTN	548	1e-53	86
1957	16	701051552H1	SOYMON032	g609224	BLASTN	744	1e-53	85
1958	16	700944049H1	SOYMON024	g609224	BLASTN	744	1e-53	85
1959	16	701120261H1	SOYMON037	g609224	BLASTN	746	1e-53	84
1960	16	700897524H1	SOYMON027	g609224	BLASTN	751	1e-53	84
1961	16	LIB3049-042-Q1-E1-F7	LIB3049	g450548	BLASTN	761	1e-53	75
1962	16	700896909H1	SOYMON027	g429105	BLASTN	536	1e-52	82
1963	16	700974820H1	SOYMON005	g1724103	BLASTN	582	1e-52	86
1964	16	701143224H1	SOYMON038	g609224	BLASTN	731	1e-52	84
1965	16	701010322H1	SOYMON019	g609224	BLASTN	731	1e-52	85
1966	16	701130510H1	SOYMON038	g609224	BLASTN	732	1e-52	83
1967	16	700724904H1	SOYMON009	g609224	BLASTN	733	1e-52	85
1968	16	701119845H1	SOYMON037	g609224	BLASTN	734	1e-52	85
1969	16	701107871H1	SOYMON036	g16508	BLASTN	737	1e-52	86
1970	16	700983380H1	SOYMON009	g609224	BLASTN	738	1e-52	84
1971	16	700896469H1	SOYMON027	g16508	BLASTN	741	1e-52	86
1972	16	700792178H1	SOYMON011	g16508	BLASTN	742	1e-52	85
1973	16	701099906H1	SOYMON028	g16508	BLASTN	431	1e-51	89
1974	16	701134724H2	SOYMON038	g16508	BLASTN	553	1e-51	85
1975	16	700749712H1	SOYMON013	g16508	BLASTN	689	1e-51	87
1976	16	700982567H1	SOYMON009	g609224	BLASTN	723	1e-51	85
1977	16	701013758H1	SOYMON019	g609224	BLASTN	723	1e-51	85
1978	16	700868508H1	SOYMON016	g609224	BLASTN	723	1e-51	85
1979	16	700749316H1	SOYMON013	g609224	BLASTN	723	1e-51	85
1980	16	701045367H1	SOYMON032	g16508	BLASTN	724	1e-51	91
1981	16	701205494H1	SOYMON035	g609224	BLASTN	724	1e-51	84
1982	16	700556784H1	SOYMON001	g16508	BLASTN	724	1e-51	91
1983	16	701099879H1	SOYMON028	g16508	BLASTN	726	1e-51	80
1984	16	701131671H1	SOYMON038	g609224	BLASTN	728	1e-51	85
1985	16	701011505H1	SOYMON019	g609224	BLASTN	728	1e-51	85
1986	16	701009973H2	SOYMON019	g609224	BLASTN	728	1e-51	85
1987	16	700793520H1	SOYMON017	g609224	BLASTN	728	1e-51	85
1988	16	700753622H1	SOYMON014	g609224	BLASTN	728	1e-51	85
1989	16	700984052H1	SOYMON009	g609224	BLASTN	728	1e-51	83
1990	16	701108521H1	SOYMON036	g609224	BLASTN	728	1e-51	85
1991	16	700957203H1	SOYMON022	g1724103	BLASTN	464	1e-50	86
1992	16	700554120H1	SOYMON001	g609224	BLASTN	528	1e-50	84
1993	16	700555954H1	SOYMON001	g167961	BLASTN	528	1e-50	86
1994	16	701124141H1	SOYMON037	g609224	BLASTN	537	1e-50	85
1995	16	701003232H1	SOYMON019	g16508	BLASTN	544	1e-50	88
1996	16	700653112H1	SOYMON003	g16508	BLASTN	559	1e-50	91
1997	16	701103353H1	SOYMON028	g1655577	BLASTN	581	1e-50	84
1998	16	700562790H1	SOYMON002	g497899	BLASTN	708	1e-50	86
1999	16	701097303H1	SOYMON028	g609224	BLASTN	711	1e-50	84
2000	16	700561195H1	SOYMON002	g609224	BLASTN	712	1e-50	84
2001	16	701121162H1	SOYMON037	g16508	BLASTN	712	1e-50	88
2002	16	700981317H1	SOYMON009	g16508	BLASTN	714	1e-50	89
2003	16	700981595H1	SOYMON009	g16508	BLASTN	716	1e-50	86
2004	16	700994305H1	SOYMON011	g609224	BLASTN	528	1e-49	85
2005	16	701101565H1	SOYMON028	g429105	BLASTN	552	1e-49	82
2006	16	701103456H1	SOYMON028	g429105	BLASTN	617	1e-49	81
2007	16	700993331H1	SOYMON011	g16508	BLASTN	622	1e-49	86
2008	16	LIB3052-011-Q1-N1-B12	LIB3052	g16508	BLASTN	695	1e-49	85
2009	16	701140613H1	SOYMON038	g609224	BLASTN	697	1e-49	85

2010	16	700745426H1	SOYMON013	g16508	BLASTN	697	1e-49	84
2011	16	701046530H1	SOYMON032	g609224	BLASTN	697	1e-49	84
2012	16	701003787H1	SOYMON019	g609224	BLASTN	697	1e-49	85
2013	16	700840830H1	SOYMON020	g1724103	BLASTN	699	1e-49	79
2014	16	700901588H1	SOYMON027	g609224	BLASTN	702	1e-49	85
2015	16	701037824H1	SOYMON029	g497899	BLASTN	702	1e-49	87
2016	16	701131888H1	SOYMON038	g609224	BLASTN	702	1e-49	85
2017	16	701009947H2	SOYMON019	g609224	BLASTN	702	1e-49	85
2018	16	700729216H1	SOYMON009	g497899	BLASTN	702	1e-49	87
2019	16	700831705H1	SOYMON019	g609224	BLASTN	702	1e-49	85
2020	16	700900329H1	SOYMON027	g429105	BLASTN	705	1e-49	88
2021	16	700563946H1	SOYMON002	g16508	BLASTN	705	1e-49	83
2022	16	700808370H1	SOYMON024	g609556	BLASTN	459	1e-48	83
2023	16	LIB3040-001-Q1-E1-H4	LIB3040	g1724103	BLASTN	475	1e-48	76
2024	16	700989482H1	SOYMON011	g16508	BLASTN	498	1e-48	86
2025	16	701001311H1	SOYMON018	g16508	BLASTN	655	1e-48	85
2026	16	700734733H1	SOYMON010	g497899	BLASTN	682	1e-48	87
2027	16	701013070H1	SOYMON019	g497899	BLASTN	682	1e-48	87
2028	16	701120989H1	SOYMON037	g16508	BLASTN	683	1e-48	91
2029	16	LIB3049-025-Q1-E1-B4	LIB3049	g167961	BLASTN	687	1e-48	82
2030	16	700808455H1	SOYMON024	g16508	BLASTN	688	1e-48	87
2031	16	701101319H1	SOYMON028	g16508	BLASTN	688	1e-48	91
2032	16	701037087H1	SOYMON029	g609224	BLASTN	689	1e-48	85
2033	16	701205225H1	SOYMON035	g16508	BLASTN	689	1e-48	84
2034	16	701136807H1	SOYMON038	g609224	BLASTN	690	1e-48	85
2035	16	701118459H1	SOYMON037	g609224	BLASTN	690	1e-48	85
2036	16	700942825H1	SOYMON024	g16508	BLASTN	691	1e-48	85
2037	16	701139796H1	SOYMON038	g497899	BLASTN	692	1e-48	87
2038	16	700557040H1	SOYMON001	g609224	BLASTN	692	1e-48	85
2039	16	701015715H1	SOYMON038	g497899	BLASTN	692	1e-48	87
2040	16	700726424H1	SOYMON009	g497899	BLASTN	693	1e-48	87
2041	16	700790811H1	SOYMON011	g497899	BLASTN	693	1e-48	87
2042	16	LIB3040-038-Q1-E1-E5	LIB3040	g16508	BLASTN	709	1e-48	85
2043	16	700896626H1	SOYMON027	g429107	BLASTN	430	1e-47	84
2044	16	700562158H1	SOYMON002	g609224	BLASTN	518	1e-47	84
2045	16	700747095H1	SOYMON013	g16508	BLASTN	538	1e-47	87
2046	16	700726532H1	SOYMON009	g609224	BLASTN	622	1e-47	85
2047	16	701137686H1	SOYMON038	g497899	BLASTN	671	1e-47	85
2048	16	701009148H1	SOYMON019	g16508	BLASTN	673	1e-47	91
2049	16	701048279H1	SOYMON032	g16508	BLASTN	674	1e-47	86
2050	16	701120185H1	SOYMON037	g16508	BLASTN	674	1e-47	91
2051	16	701040355H1	SOYMON029	g497899	BLASTN	675	1e-47	86
2052	16	701130203H1	SOYMON037	g609224	BLASTN	678	1e-47	83
2053	16	701015794H1	SOYMON038	g16508	BLASTN	681	1e-47	91
2054	16	700732181H1	SOYMON010	g16508	BLASTN	681	1e-47	86
2055	16	LIB3050-018-Q1-E1-D10	LIB3050	g2305013	BLASTN	696	1e-47	81
2056	16	700899157H1	SOYMON027	g609224	BLASTN	486	1e-46	84
2057	16	701119905H1	SOYMON037	g497899	BLASTN	507	1e-46	86
2058	16	701062873H1	SOYMON033	g497899	BLASTN	509	1e-46	88
2059	16	701210886H1	SOYMON035	g16508	BLASTN	540	1e-46	84
2060	16	700893278H1	SOYMON024	g16508	BLASTN	542	1e-46	77
2061	16	701036970H1	SOYMON029	g16508	BLASTN	579	1e-46	88
2062	16	700901932H1	SOYMON027	g609224	BLASTN	640	1e-46	85
2063	16	701044214H1	SOYMON032	g429105	BLASTN	658	1e-46	85

2064	16	701056917H1	SOYMON033	g16508	BLASTN	661	1e-46	85
2065	16	700981560H1	SOYMON009	g497899	BLASTN	661	1e-46	87
2066	16	701213107H1	SOYMON035	g609224	BLASTN	661	1e-46	83
2067	16	700834235H1	SOYMON019	g497899	BLASTN	661	1e-46	87
2068	16	700749413H1	SOYMON013	g16508	BLASTN	662	1e-46	85
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2071	16	701206618H1	SOYMON035	g16508	BLASTN	663	1e-46	91
2072	16	701131927H1	SOYMON038	g497899	BLASTN	666	1e-46	87
2073	16	701119069H1	SOYMON037	g497899	BLASTN	666	1e-46	87
2074	16	701106908H1	SOYMON036	g497899	BLASTN	666	1e-46	87
2075	16	701103063H1	SOYMON028	g497899	BLASTN	666	1e-46	87
2076	16	700686659H1	SOYMON008	g166873	BLASTN	667	1e-46	86
2077	16	701040620H1	SOYMON029	g16508	BLASTN	669	1e-46	86
2078	16	701012134H1	SOYMON019	g497899	BLASTN	669	1e-46	85
2079	16	700957508H1	SOYMON022	g1724103	BLASTN	669	1e-46	82
2080	16	701008548H1	SOYMON019	g497899	BLASTN	500	1e-45	87
2081	16	700978303H1	SOYMON009	g497899	BLASTN	506	1e-45	87
2082	16	701119990H1	SOYMON037	g497899	BLASTN	512	1e-45	87
2083	16	700646222H1	SOYMON012	g16508	BLASTN	526	1e-45	85
2084	16	701040631H1	SOYMON029	g609556	BLASTN	542	1e-45	87
2085	16	700894009H1	SOYMON024	g497899	BLASTN	646	1e-45	87
2086	16	700908702H1	SOYMON022	g609224	BLASTN	646	1e-45	85
2087	16	700838642H1	SOYMON020	g609224	BLASTN	646	1e-45	85
2088	16	700832089H1	SOYMON019	g609224	BLASTN	646	1e-45	85
2089	16	700978279H1	SOYMON009	g609224	BLASTN	647	1e-45	80
2090	16	700555058H1	SOYMON001	g16508	BLASTN	649	1e-45	84
2091	16	700906478H1	SOYMON022	g497899	BLASTN	651	1e-45	87
2092	16	700897237H1	SOYMON027	g16508	BLASTN	653	1e-45	86
2093	16	701125786H1	SOYMON037	g16508	BLASTN	654	1e-45	84
2094	16	700562836H1	SOYMON002	g16508	BLASTN	655	1e-45	84
2095	16	700665969H1	SOYMON005	g497899	BLASTN	656	1e-45	87
2096	16	701009838H1	SOYMON019	g497899	BLASTN	656	1e-45	87
2097	16	701211770H1	SOYMON035	g16508	BLASTN	657	1e-45	84
2098	16	700654550H1	SOYMON004	g16508	BLASTN	657	1e-45	83
2099	16	700567949H1	SOYMON002	g16508	BLASTN	657	1e-45	84
2100	16	701048175H1	SOYMON032	g16508	BLASTN	658	1e-45	91
2101	16	700901037H1	SOYMON027	g16508	BLASTN	658	1e-45	86
2102	16	700656917H1	SOYMON004	g1724103	BLASTN	407	1e-44	81
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2104	16	701002808H1	SOYMON019	g497899	BLASTN	503	1e-44	87
2105	16	701107115H1	SOYMON036	g497899	BLASTN	507	1e-44	87
2106	16	701122669H1	SOYMON037	g16508	BLASTN	636	1e-44	82
2107	16	700894409H1	SOYMON024	g497899	BLASTN	636	1e-44	87
2108	16	700848374H1	SOYMON021	g497899	BLASTN	636	1e-44	87
2109	16	700902420H1	SOYMON027	g497899	BLASTN	636	1e-44	87
2110	16	700747450H1	SOYMON013	g16508	BLASTN	638	1e-44	91
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2112	16	700731302H1	SOYMON010	g16508	BLASTN	639	1e-44	85
2113	16	701099068H1	SOYMON028	g2305013	BLASTN	640	1e-44	81
2114	16	700975556H1	SOYMON009	g450548	BLASTN	641	1e-44	84
2115	16	700889660H1	SOYMON024	g16508	BLASTN	642	1e-44	86
2116	16	700951744H1	SOYMON022	g16508	BLASTN	642	1e-44	86
2117	16	700750251H1	SOYMON013	g609224	BLASTN	643	1e-44	86

2118	16	700834611H1	SOYMON019	g16508	BLASTN	643	1e-44	91
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2120	16	700673803H1	SOYMON007	g16508	BLASTN	644	1e-44	84
2121	16	700958721H1	SOYMON022	g16508	BLASTN	645	1e-44	86
2122	16	700985717H1	SOYMON009	g16508	BLASTN	646	1e-44	82
2123	16	LIB3049-032-Q1-E1-A5	LIB3049	g16508	BLASTN	662	1e-44	76
2124	16	700733454H1	SOYMON010	g609224	BLASTN	432	1e-43	85
2125	16	700987656H1	SOYMON009	g497899	BLASTN	476	1e-43	87
2126	16	700755957H1	SOYMON014	g497899	BLASTN	483	1e-43	86
2127	16	700749331H1	SOYMON013	g2305013	BLASTN	521	1e-43	84
2128	16	700830984H1	SOYMON019	g497899	BLASTN	622	1e-43	86
2129	16	700906176H1	SOYMON022	g16508	BLASTN	623	1e-43	86
2130	16	701100070H2	SOYMON028	g16508	BLASTN	623	1e-43	91
2131	16	700954053H1	SOYMON022	g497899	BLASTN	624	1e-43	88
2132	16	700833949H1	SOYMON019	g497899	BLASTN	624	1e-43	88
2133	16	700976710H1	SOYMON009	g2305013	BLASTN	625	1e-43	79
2134	16	701117925H2	SOYMON037	g16508	BLASTN	625	1e-43	84
2135	16	700748825H1	SOYMON013	g16508	BLASTN	626	1e-43	84
2136	16	700899651H1	SOYMON027	g16508	BLASTN	626	1e-43	84
2137	16	700746739H1	SOYMON013	g497899	BLASTN	626	1e-43	88
2138	16	701009051H1	SOYMON019	g16508	BLASTN	628	1e-43	91
2139	16	700754423H1	SOYMON014	g16508	BLASTN	628	1e-43	91
2140	16	700760701H1	SOYMON015	g16508	BLASTN	628	1e-43	83
2141	16	701098124H1	SOYMON028	g2305013	BLASTN	628	1e-43	78
2142	16	700984979H1	SOYMON009	g16508	BLASTN	628	1e-43	91
2143	16	700565262H1	SOYMON002	g497899	BLASTN	629	1e-43	86
2144	16	700954979H1	SOYMON022	g497899	BLASTN	630	1e-43	87
2145	16	700834601H1	SOYMON019	g497899	BLASTN	630	1e-43	87
2146	16	700953156H1	SOYMON022	g609224	BLASTN	630	1e-43	85
2147	16	700865217H1	SOYMON016	g16508	BLASTN	631	1e-43	77
2148	16	700852947H1	SOYMON023	g16508	BLASTN	631	1e-43	84
2149	16	700943547H1	SOYMON024	g497899	BLASTN	631	1e-43	86
2150	16	701139277H1	SOYMON038	g609224	BLASTN	632	1e-43	82
2151	16	700900155H1	SOYMON027	g429105	BLASTN	264	1e-42	81
2152	16	700970581H1	SOYMON005	g497899	BLASTN	296	1e-42	86
2153	16	700654232H1	SOYMON003	g497899	BLASTN	348	1e-42	88
2154	16	700743608H1	SOYMON012	g497899	BLASTN	354	1e-42	86
2155	16	700986620H1	SOYMON009	g609224	BLASTN	360	1e-42	84
2156	16	700562590H1	SOYMON002	g16508	BLASTN	476	1e-42	83
2157	16	701119690H1	SOYMON037	g167961	BLASTN	506	1e-42	84
2158	16	700951730H1	SOYMON022	g497899	BLASTN	507	1e-42	87
2159	16	701004525H1	SOYMON019	g16508	BLASTN	516	1e-42	91
2160	16	700892637H1	SOYMON024	g16508	BLASTN	523	1e-42	81
2161	16	700560679H1	SOYMON001	g16508	BLASTN	526	1e-42	83
2162	16	700897888H1	SOYMON027	g16508	BLASTN	611	1e-42	86
2163	16	700547973H1	SOYMON001	g497899	BLASTN	611	1e-42	84
2164	16	701012166H1	SOYMON019	g497899	BLASTN	612	1e-42	87
2165	16	700747550H1	SOYMON013	g16508	BLASTN	613	1e-42	84
2166	16	700946136H1	SOYMON024	g16508	BLASTN	615	1e-42	85
2167	16	700665932H1	SOYMON005	g16508	BLASTN	616	1e-42	84
2168	16	701010969H1	SOYMON019	g16508	BLASTN	618	1e-42	84
2169	16	700962515H1	SOYMON022	g16508	BLASTN	618	1e-42	91
2170	16	700895685H1	SOYMON027	g429107	BLASTN	618	1e-42	77
2171	16	700752015H1	SOYMON014	g16508	BLASTN	620	1e-42	84

2172	16	701015704H1	SOYMON038	g16508	BLASTN	620	1e-42	85
2173	16	700966710H1	SOYMON028	g16508	BLASTN	620	1e-42	83
2174	16	701004268H1	SOYMON019	g16508	BLASTN	620	1e-42	86
2175	16	701138901H1	SOYMON038	g497899	BLASTN	285	1e-41	85
2176	16	700648891H1	SOYMON003	g16508	BLASTN	481	1e-41	84
2177	16	700741421H1	SOYMON012	g609224	BLASTN	511	1e-41	82
2178	16	700555653H1	SOYMON001	g166873	BLASTN	512	1e-41	85
2179	16	700958314H1	SOYMON022	g497899	BLASTN	599	1e-41	87
2180	16	700960055H1	SOYMON022	g497899	BLASTN	599	1e-41	87
2181	16	700901925H1	SOYMON027	g16508	BLASTN	599	1e-41	84
2182	16	700741987H1	SOYMON012	g497899	BLASTN	599	1e-41	87
2183	16	700662837H1	SOYMON005	g497899	BLASTN	599	1e-41	87
2184	16	701060927H1	SOYMON033	g497899	BLASTN	599	1e-41	87
2185	16	700562973H1	SOYMON002	g16508	BLASTN	601	1e-41	87
2186	16	700754586H1	SOYMON014	g16508	BLASTN	603	1e-41	84
2187	16	701132674H1	SOYMON038	g16508	BLASTN	604	1e-41	85
2188	16	701010012H2	SOYMON019	g609224	BLASTN	604	1e-41	84
2189	16	701123643H1	SOYMON037	g16508	BLASTN	604	1e-41	85
2190	16	700894196H1	SOYMON024	g16508	BLASTN	604	1e-41	85
2191	16	700899248H1	SOYMON027	g16508	BLASTN	604	1e-41	85
2192	16	701207680H1	SOYMON035	g16508	BLASTN	605	1e-41	84
2193	16	700899210H1	SOYMON027	g16508	BLASTN	605	1e-41	84
2194	16	700898762H1	SOYMON027	g16508	BLASTN	605	1e-41	87
2195	16	700983636H1	SOYMON009	g497899	BLASTN	606	1e-41	87
2196	16	700845511H1	SOYMON021	g16508	BLASTN	606	1e-41	89
2197	16	700563416H1	SOYMON002	g16508	BLASTN	606	1e-41	83
2198	16	701004041H1	SOYMON019	g16508	BLASTN	607	1e-41	84
2199	16	701097184H1	SOYMON028	g16508	BLASTN	607	1e-41	87
2200	16	700750691H1	SOYMON014	g609224	BLASTN	608	1e-41	84
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2202	16	700973888H1	SOYMON005	g609224	BLASTN	609	1e-41	85
2203	16	700831972H1	SOYMON019	g16508	BLASTN	609	1e-41	85
2204	16	701010090H2	SOYMON019	g16508	BLASTN	610	1e-41	87
2205	16	700890067H1	SOYMON024	g16508	BLASTN	610	1e-41	87
2206	16	700833227H1	SOYMON019	g16508	BLASTN	610	1e-41	85
2207	16	700563171H1	SOYMON002	g16508	BLASTN	370	1e-40	90
2208	16	701121457H1	SOYMON037	g609224	BLASTN	372	1e-40	85
2209	16	700566851H1	SOYMON002	g16508	BLASTN	402	1e-40	82
2210	16	700753221H1	SOYMON014	g609224	BLASTN	422	1e-40	84
2211	16	700749039H1	SOYMON013	g16508	BLASTN	475	1e-40	84
2212	16	700558940H1	SOYMON001	g16508	BLASTN	493	1e-40	80
2213	16	700757695H1	SOYMON015	g16508	BLASTN	526	1e-40	83
2214	16	701102262H1	SOYMON028	g2305013	BLASTN	587	1e-40	81
2215	16	700747484H1	SOYMON013	g16508	BLASTN	587	1e-40	85
2216	16	700891875H1	SOYMON024	g16508	BLASTN	587	1e-40	85
2217	16	700894439H1	SOYMON024	g497899	BLASTN	588	1e-40	88
2218	16	701062577H1	SOYMON033	g16508	BLASTN	588	1e-40	87
2219	16	700964366H1	SOYMON022	g16508	BLASTN	589	1e-40	85
2220	16	701210016H1	SOYMON035	g16508	BLASTN	589	1e-40	83
2221	16	700901573H1	SOYMON027	g16508	BLASTN	589	1e-40	85
2222	16	700750533H1	SOYMON014	g16508	BLASTN	589	1e-40	85
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2224	16	700891965H1	SOYMON024	g497899	BLASTN	589	1e-40	87
2225	16	700865001H1	SOYMON016	g16508	BLASTN	589	1e-40	85

2226	16	700654923H1	SOYMON004	g16508	BLASTN	590	1e-40	84
2227	16	700746169H1	SOYMON013	g16508	BLASTN	590	1e-40	84
2228	16	700745792H1	SOYMON013	g16508	BLASTN	591	1e-40	85
2229	16	700966953H1	SOYMON029	g16508	BLASTN	591	1e-40	85
2230	16	701141657H1	SOYMON038	g16508	BLASTN	592	1e-40	86
2231	16	700943078H1	SOYMON024	g16508	BLASTN	598	1e-40	85
2232	16	701137073H1	SOYMON038	g16508	BLASTN	598	1e-40	90
2233	16	700966730H1	SOYMON028	g16508	BLASTN	598	1e-40	84
2234	16	701137731H1	SOYMON038	g16508	BLASTN	371	1e-39	84
2235	16	701044050H1	SOYMON032	g497899	BLASTN	428	1e-39	89
2236	16	700993309H1	SOYMON011	g497899	BLASTN	434	1e-39	88
2237	16	700874483H1	SOYMON018	g497899	BLASTN	434	1e-39	87
2238	16	701014866H1	SOYMON019	g16508	BLASTN	454	1e-39	89
2239	16	700894603H1	SOYMON024	g16508	BLASTN	454	1e-39	85
2240	16	700958677H1	SOYMON022	g497899	BLASTN	455	1e-39	87
2241	16	700754001H1	SOYMON014	g609224	BLASTN	518	1e-39	85
2242	16	700946430H1	SOYMON024	g497899	BLASTN	578	1e-39	87
2243	16	700729892H1	SOYMON009	g497899	BLASTN	578	1e-39	87
2244	16	700753230H1	SOYMON014	g497899	BLASTN	578	1e-39	87
2245	16	700562584H1	SOYMON002	g16508	BLASTN	578	1e-39	82
2246	16	700842880H1	SOYMON020	g16508	BLASTN	580	1e-39	82
2247	16	700852561H1	SOYMON023	g16508	BLASTN	582	1e-39	85
2248	16	700941290H1	SOYMON024	g497899	BLASTN	583	1e-39	87
2249	16	700750361H1	SOYMON013	g497899	BLASTN	584	1e-39	80
2250	16	700841915H1	SOYMON020	g16508	BLASTN	585	1e-39	85
2251	16	700944462H1	SOYMON024	g16508	BLASTN	585	1e-39	85
2252	16	700548174H1	SOYMON002	g16508	BLASTN	342	1e-38	82
2253	16	700942324H1	SOYMON024	g497899	BLASTN	403	1e-38	87
2254	16	700788450H1	SOYMON011	g497899	BLASTN	416	1e-38	86
2255	16	701127954H1	SOYMON037	g16508	BLASTN	449	1e-38	76
2256	16	700742223H1	SOYMON012	g1724103	BLASTN	451	1e-38	77
2257	16	701102722H1	SOYMON028	g16508	BLASTN	466	1e-38	92
2258	16	701046957H1	SOYMON032	g16508	BLASTN	517	1e-38	85
2259	16	700836169H1	SOYMON019	g497899	BLASTN	517	1e-38	87
2260	16	700697967H1	SOYMON015	g16508	BLASTN	521	1e-38	84
2261	16	700678867H1	SOYMON007	g609224	BLASTN	566	1e-38	82
2262	16	700838753H1	SOYMON020	g16508	BLASTN	567	1e-38	85
2263	16	700726468H1	SOYMON009	g16508	BLASTN	570	1e-38	85
2264	16	701014631H1	SOYMON019	g497899	BLASTN	571	1e-38	87
2265	16	700728719H1	SOYMON009	g497899	BLASTN	573	1e-38	87
2266	16	700962582H1	SOYMON022	g497899	BLASTN	573	1e-38	87
2267	16	701123183H1	SOYMON037	g16508	BLASTN	573	1e-38	84
2268	16	700752347H1	SOYMON014	g16508	BLASTN	574	1e-38	87
2269	16	701103315H1	SOYMON028	g16508	BLASTN	394	1e-37	86
2270	16	700750955H1	SOYMON014	g497899	BLASTN	414	1e-37	88
2271	16	700867493H1	SOYMON016	g609224	BLASTN	433	1e-37	85
2272	16	701054760H1	SOYMON032	g16508	BLASTN	475	1e-37	85
2273	16	700900903H1	SOYMON027	g16508	BLASTN	501	1e-37	84
2274	16	701102294H1	SOYMON028	g16508	BLASTN	502	1e-37	87
2275	16	701124034H1	SOYMON037	g16508	BLASTN	512	1e-37	82
2276	16	700981476H1	SOYMON009	g16508	BLASTN	526	1e-37	83
2277	16	700903712H1	SOYMON022	g609224	BLASTN	552	1e-37	85
2278	16	700648751H1	SOYMON003	g1655577	BLASTN	554	1e-37	86
2279	16	700829930H1	SOYMON019	g609224	BLASTN	556	1e-37	84



2280	16	700758078H1	SOYMON015	g16508	BLASTN	556	1e-37	84
2281	16	700754248H1	SOYMON014	g16508	BLASTN	557	1e-37	81
2282	16	700834650H1	SOYMON019	g609224	BLASTN	557	1e-37	85
2283	16	700746279H1	SOYMON013	g16508	BLASTN	558	1e-37	82
2284	16	700869124H1	SOYMON016	g16508	BLASTN	559	1e-37	83
2285	16	701100219H1	SOYMON028	g2305013	BLASTN	560	1e-37	80
2286	16	700992589H1	SOYMON011	g497899	BLASTN	560	1e-37	80
2287	16	700865642H1	SOYMON016	g16508	BLASTN	562	1e-37	84
2288	16	701068835H1	SOYMON034	g429105	BLASTN	321	1e-36	86
2289	16	700746908H1	SOYMON013	g16508	BLASTN	373	1e-36	81
2290	16	700954719H1	SOYMON022	g16508	BLASTN	475	1e-36	86
2291	16	701042790H1	SOYMON029	g497899	BLASTN	489	1e-36	88
2292	16	700835259H1	SOYMON019	g497899	BLASTN	507	1e-36	88
2293	16	700565816H1	SOYMON002	g16508	BLASTN	519	1e-36	82
2294	16	700895811H1	SOYMON027	g609224	BLASTN	540	1e-36	83
2295	16	700975709H1	SOYMON009	g609224	BLASTN	541	1e-36	84
2296	16	700751645H1	SOYMON014	g16508	BLASTN	542	1e-36	85
2297	16	700567087H1	SOYMON002	g16508	BLASTN	542	1e-36	85
2298	16	700893027H1	SOYMON024	g609224	BLASTN	542	1e-36	85
2299	16	700891185H1	SOYMON024	g609224	BLASTN	543	1e-36	84
2300	16	700851376H1	SOYMON023	g16508	BLASTN	543	1e-36	83
2301	16	700898322H1	SOYMON027	g609224	BLASTN	547	1e-36	85
2302	16	700762881H1	SOYMON015	g16508	BLASTN	323	1e-35	85
2303	16	700960023H1	SOYMON022	g16508	BLASTN	370	1e-35	90
2304	16	701136486H1	SOYMON038	g497899	BLASTN	378	1e-35	85
2305	16	700987688H1	SOYMON009	g497899	BLASTN	387	1e-35	83
2306	16	700836508H1	SOYMON020	g16508	BLASTN	431	1e-35	85
2307	16	700990779H1	SOYMON011	g167961	BLASTN	471	1e-35	86
2308	16	700753019H1	SOYMON014	g497899	BLASTN	476	1e-35	83
2309	16	701138490H1	SOYMON038	g16508	BLASTN	481	1e-35	83
2310	16	700751771H1	SOYMON014	g609224	BLASTN	528	1e-35	84
2311	16	701048324H1	SOYMON032	g609224	BLASTN	529	1e-35	84
2312	16	700835739H1	SOYMON019	g16508	BLASTN	532	1e-35	85
2313	16	700986295H1	SOYMON009	g497899	BLASTN	533	1e-35	86
2314	16	701205693H1	SOYMON035	g16508	BLASTN	535	1e-35	84
2315	16	700865674H1	SOYMON016	g609224	BLASTN	536	1e-35	85
2316	16	700943777H1	SOYMON024	g16508	BLASTN	536	1e-35	84
2317	16	700968207H1	SOYMON035	g609224	BLASTN	536	1e-35	85
2318	16	700898370H1	SOYMON027	g609224	BLASTN	536	1e-35	85
2319	16	700987890H1	SOYMON009	g609224	BLASTN	536	1e-35	85
2320	16	700896016H1	SOYMON027	g609224	BLASTN	536	1e-35	85
2321	16	700554440H1	SOYMON001	g16508	BLASTN	283	1e-34	82
2322	16	700789855H2	SOYMON011	g16508	BLASTN	293	1e-34	91
2323	16	701101393H1	SOYMON028	g16508	BLASTN	311	1e-34	84
2324	16	700946335H1	SOYMON024	g16508	BLASTN	323	1e-34	86
2325	16	701207420H1	SOYMON035	g169664	BLASTN	350	1e-34	88
2326	16	700733034H1	SOYMON010	g16508	BLASTN	384	1e-34	81
2327	16	700790653H2	SOYMON011	g497899	BLASTN	399	1e-34	81
2328	16	700901808H1	SOYMON027	g497899	BLASTN	430	1e-34	81
2329	16	700760954H1	SOYMON015	g609224	BLASTN	514	1e-34	84
2330	16	700896206H1	SOYMON027	g16508	BLASTN	516	1e-34	85
2331	16	701039157H1	SOYMON029	g609224	BLASTN	520	1e-34	76
2332	16	700991056H1	SOYMON011	g166873	BLASTN	520	1e-34	80
2333	16	701097173H1	SOYMON028	g609224	BLASTN	521	1e-34	85

2334	16	700900989H1	SOYMON027	g497899	BLASTN	521	1e-34	86
2335	16	700895584H1	SOYMON027	g609224	BLASTN	521	1e-34	85
2336	16	700726668H1	SOYMON009	g16508	BLASTN	521	1e-34	85
2337	16	701011191H1	SOYMON019	g16508	BLASTN	521	1e-34	85
2338	16	701100827H1	SOYMON028	g16508	BLASTN	522	1e-34	84
2339	16	701156732H1	SOYMON031	g16508	BLASTN	522	1e-34	84
2340	16	700889752H1	SOYMON024	g609224	BLASTN	523	1e-34	84
2341	16	701105873H1	SOYMON036	g16508	BLASTN	524	1e-34	87
2342	16	700958346H1	SOYMON022	g497899	BLASTN	525	1e-34	84
2343	16	700962972H1	SOYMON022	g609224	BLASTN	525	1e-34	84
2344	16	701055918H1	SOYMON032	g16508	BLASTN	526	1e-34	85
2345	16	701123061H1	SOYMON037	g16508	BLASTN	526	1e-34	85
2346	16	701109796H1	SOYMON036	g166873	BLASTN	335	1e-33	86
2347	16	701207350H1	SOYMON035	g497899	BLASTN	349	1e-33	89
2348	16	700835692H1	SOYMON019	g16508	BLASTN	377	1e-33	81
2349	16	700729083H1	SOYMON009	g16508	BLASTN	448	1e-33	85
2350	16	LIB3040-049-Q1-E1-E8	LIB3040	g16508	BLASTN	464	1e-33	80
2351	16	700896567H1	SOYMON027	g497899	BLASTN	502	1e-33	81
2352	16	700668032H1	SOYMON006	g609224	BLASTN	504	1e-33	85
2353	16	700895062H1	SOYMON024	g609224	BLASTN	505	1e-33	83
2354	16	700961178H1	SOYMON022	g16508	BLASTN	506	1e-33	84
2355	16	701202590H1	SOYMON035	g497899	BLASTN	508	1e-33	86
2356	16	700754960H1	SOYMON014	g16508	BLASTN	512	1e-33	83
2357	16	700748584H1	SOYMON013	g16508	BLASTN	512	1e-33	84
2358	16	700791669H1	SOYMON011	g609224	BLASTN	512	1e-33	84
2359	16	701210405H1	SOYMON035	g16508	BLASTN	526	1e-33	83
2360	16	LIB3040-054-Q1-E1-D8	LIB3040	g16508	BLASTN	531	1e-33	80
2361	16	700791176H1	SOYMON011	g497899	BLASTN	311	1e-32	87
2362	16	LIB3049-029-Q1-E1-C5	LIB3049	g167961	BLASTN	368	1e-32	73
2363	16	700893458H1	SOYMON024	g497899	BLASTN	380	1e-32	87
2364	16	700829728H1	SOYMON019	g497899	BLASTN	490	1e-32	88
2365	16	700833137H1	SOYMON019	g609224	BLASTN	490	1e-32	85
2366	16	701204592H2	SOYMON035	g609224	BLASTN	493	1e-32	86
2367	16	700834821H1	SOYMON019	g609224	BLASTN	494	1e-32	78
2368	16	700757167H1	SOYMON015	g16508	BLASTN	494	1e-32	85
2369	16	701203730H2	SOYMON035	g16508	BLASTN	496	1e-32	82
2370	16	701044359H1	SOYMON032	g609224	BLASTN	497	1e-32	84
2371	16	701010095H2	SOYMON019	g16508	BLASTN	498	1e-32	86
2372	16	701100059H2	SOYMON028	g16508	BLASTN	499	1e-32	80
2373	16	700844256H1	SOYMON021	g16508	BLASTN	500	1e-32	80
2374	16	700968296H1	SOYMON035	g497899	BLASTN	501	1e-32	85
2375	16	701110404H1	SOYMON036	g16508	BLASTN	333	1e-31	84
2376	16	700869025H1	SOYMON016	g497899	BLASTN	340	1e-31	86
2377	16	700556245H1	SOYMON001	g609224	BLASTN	355	1e-31	84
2378	16	700566166H1	SOYMON002	g726031	BLASTN	402	1e-31	77
2379	16	700941481H1	SOYMON024	g497899	BLASTN	423	1e-31	79
2380	16	700896940H1	SOYMON027	g169664	BLASTN	477	1e-31	85
2381	16	701145429H1	SOYMON031	g609224	BLASTN	478	1e-31	85
2382	16	700946496H1	SOYMON024	g609224	BLASTN	488	1e-31	86
2383	16	700902149H1	SOYMON027	g1655577	BLASTN	488	1e-31	87
2384	16	700653072H1	SOYMON003	g497899	BLASTN	489	1e-31	88
2385	16	701000261H1	SOYMON018	g16508	BLASTN	495	1e-31	78
2386	16	701039358H1	SOYMON029	g609224	BLASTN	320	1e-30	85
2387	16	701156951H1	SOYMON031	g609224	BLASTN	467	1e-30	86

2388	16	701040880H1	SOYMON029	g497899	BLASTN	477	1e-30	89
2389	16	700979632H2	SOYMON009	g609224	BLASTN	490	1e-30	83
2390	16	700983678H1	SOYMON009	g16508	BLASTN	192	1e-29	86
2391	16	700893644H1	SOYMON024	g609224	BLASTN	320	1e-29	85
2392	16	700962437H1	SOYMON022	g609224	BLASTN	364	1e-29	83
2393	16	701150545H1	SOYMON031	g16508	BLASTN	457	1e-29	86
2394	16	700555403H1	SOYMON001	g16508	BLASTN	458	1e-29	90
2395	16	700794307H1	SOYMON017	g16508	BLASTN	460	1e-29	87
2396	16	701137971H1	SOYMON038	g16508	BLASTN	460	1e-29	84
2397	16	700893774H1	SOYMON024	g609224	BLASTN	461	1e-29	86
2398	16	700565272H1	SOYMON002	g609224	BLASTN	464	1e-29	74
2399	16	700667161H1	SOYMON006	g16508	BLASTN	468	1e-29	75
2400	16	700752952H1	SOYMON014	g609224	BLASTN	279	1e-28	78
2401	16	700901534H1	SOYMON027	g2305013	BLASTN	288	1e-28	85
2402	16	701006617H1	SOYMON019	g497899	BLASTN	358	1e-28	79
2403	16	700753459H1	SOYMON014	g609224	BLASTN	446	1e-28	81
2404	16	701157089H1	SOYMON031	g609224	BLASTN	448	1e-28	84
2405	16	700831271H1	SOYMON019	g16508	BLASTN	453	1e-28	87
2406	16	701207325H1	SOYMON035	g167961	BLASTN	251	1e-27	81
2407	16	700566121H1	SOYMON002	g16508	BLASTN	259	1e-27	75
2408	16	701137878H1	SOYMON038	g609224	BLASTN	280	1e-27	85
2409	16	700763957H1	SOYMON019	g497899	BLASTN	430	1e-27	88
2410	16	701015858H1	SOYMON038	g609224	BLASTN	431	1e-27	86
2411	16	700742849H1	SOYMON012	g497899	BLASTN	434	1e-27	88
2412	16	701102648H1	SOYMON028	g16508	BLASTN	436	1e-27	89
2413	16	700940955H1	SOYMON024	g609556	BLASTN	438	1e-27	85
2414	16	700889805H1	SOYMON024	g16508	BLASTN	441	1e-27	89
2415	16	701061930H1	SOYMON033	g16508	BLASTN	315	1e-26	77
2416	16	700901375H1	SOYMON027	g609224	BLASTN	323	1e-26	84
2417	16	700989056H1	SOYMON011	g609224	BLASTN	345	1e-26	82
2418	16	700665988H1	SOYMON005	g960356	BLASTN	420	1e-26	79
2419	16	700893352H1	SOYMON024	g497899	BLASTN	428	1e-26	87
2420	16	701122945H1	SOYMON037	g16508	BLASTN	269	1e-25	76
2421	16	700992929H1	SOYMON011	g16508	BLASTN	328	1e-25	78
2422	16	700557564H1	SOYMON001	g609224	BLASTN	413	1e-25	85
2423	16	701102620H1	SOYMON028	g609224	BLASTN	414	1e-25	82
2424	16	701062321H1	SOYMON033	g497899	BLASTN	267	1e-24	87
2425	16	701062258H1	SOYMON033	g450548	BLASTN	391	1e-24	83
2426	16	701145703H1	SOYMON031	g609224	BLASTN	411	1e-24	85
2427	16	701144943H1	SOYMON031	g450548	BLASTN	421	1e-24	85
2428	16	701134190H1	SOYMON038	g2305013	BLASTN	268	1e-23	87
2429	16	701132116H1	SOYMON038	g450548	BLASTN	385	1e-23	85
2430	16	701144522H1	SOYMON031	g450548	BLASTN	406	1e-23	85
2431	16	700738002H1	SOYMON012	g1655578	BLASTX	205	1e-21	84
2432	16	700756686H1	SOYMON014	g16508	BLASTN	235	1e-21	76
2433	16	700649058H1	SOYMON003	g16508	BLASTN	360	1e-21	85
2434	16	701152050H1	SOYMON031	g497899	BLASTN	233	1e-20	87
2435	16	701156991H1	SOYMON031	g497900	BLASTX	146	1e-19	98
2436	16	700761949H1	SOYMON015	g497900	BLASTX	173	1e-19	98
2437	16	700988163H1	SOYMON009	g609224	BLASTN	243	1e-19	81
2438	16	700666050H1	SOYMON005	g960357	BLASTX	183	1e-18	89
2439	16	700896896H1	SOYMON027	g497900	BLASTX	95	1e-17	82
2440	16	701100249H1	SOYMON028	g497900	BLASTX	104	1e-17	79
2441	16	701061719H1	SOYMON033	g609224	BLASTN	310	1e-17	67

2442	16	700674836H1	SOYMON007	g16508	BLASTN	331	1e-17	84
2443	16	700908822H1	SOYMON022	g169665	BLASTX	171	1e-16	100
2444	16	700898670H1	SOYMON027	g166872	BLASTX	172	1e-16	93
2445	16	700650967H1	SOYMON003	g16845	BLASTX	147	1e-15	100
2446	16	701132185H1	SOYMON038	g16845	BLASTX	151	1e-15	100
2447	16	700648929H1	SOYMON003	g1033190	BLASTX	156	1e-14	96
2448	16	700960809H1	SOYMON022	g16508	BLASTN	168	1e-14	88
2449	16	701101753H1	SOYMON028	g16845	BLASTX	141	1e-13	88
2450	16	700742010H1	SOYMON012	g609224	BLASTN	273	1e-13	88
2451	16	700735080H1	SOYMON010	g166874	BLASTX	121	1e-12	86
2452	16	700979242H1	SOYMON009	g609224	BLASTN	165	1e-12	81
2453	16	700740838H1	SOYMON012	g609224	BLASTN	253	1e-12	88
2454	16	700832383H1	SOYMON019	g16845	BLASTX	82	1e-11	71
2455	16	700830938H1	SOYMON019	g16845	BLASTX	92	1e-11	90
2456	16	700564686H1	SOYMON002	g1655578	BLASTX	95	1e-11	97
2457	16	700753178H1	SOYMON014	g16845	BLASTX	107	1e-10	78
2458	16	700563834H1	SOYMON002	g497900	BLASTX	124	1e-10	100
2459	16	700897739H1	SOYMON027	g16845	BLASTX	125	1e-10	86
2460	16	700725722H1	SOYMON009	g609225	BLASTX	125	1e-10	92
2461	16	701210357H1	SOYMON035	g497900	BLASTX	128	1e-10	100
2462	16	700833654H1	SOYMON019	g609224	BLASTN	243	1e-10	87
2463	16	701010334H1	SOYMON019	g609224	BLASTN	258	1e-10	86
2464	16	700567622H1	SOYMON002	g166874	BLASTX	93	1e-9	100
2465	16	700647933H1	SOYMON003	g609225	BLASTX	120	1e-9	81
2466	16	700742255H1	SOYMON012	g429107	BLASTN	160	1e-9	86
2467	16	700981506H1	SOYMON009	g609224	BLASTN	248	1e-9	88
2468	16	700962044H1	SOYMON022	g497900	BLASTX	113	1e-8	100
2469	16	701039604H1	SOYMON029	g497900	BLASTX	113	1e-8	100
2470	16	701009941H2	SOYMON019	g609225	BLASTX	114	1e-8	70
2471	16	700976956H1	SOYMON009	g609225	BLASTX	118	1e-8	96
2472	18138	701120722H1	SOYMON037	g169664	BLASTN	514	1e-33	92
2473	18138	700946044H1	SOYMON024	g169664	BLASTN	437	1e-26	90
2474	18138	700664443H1	SOYMON005	g17262	BLASTX	162	1e-16	88
2475	18138	700665162H1	SOYMON005	g17262	BLASTX	162	1e-16	88
2476	18138	701143667H1	SOYMON038	g16961	BLASTX	123	1e-11	95
2477	18138	701099150H1	SOYMON028	g16961	BLASTX	112	1e-9	90
2478	27686	700909141H1	SOYMON022	g726027	BLASTN	723	1e-51	84
2479	27686	701145458H1	SOYMON031	g726027	BLASTN	694	1e-49	86

# ADENOSYLMETHIONINE DECARBOXYLASE ( EC 4.1.1.50 )

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
430	-700151703	700151703H1	SATMON007	g1532072	BLASTN	717	1e-50	91
431	-700165608	700165608H1	SATMON013	g1532072	BLASTN	605	1e-41	75
432	-700166868	700166868H1	SATMON013	g1532072	BLASTN	593	1e-40	90
433	-700242148	700242148H1	SATMON010	g1532048	BLASTX	142	1e-12	88
434	-700354923	700354923H1	SATMON024	g1532072	BLASTN	887	1e-74	89
435	-700422279	700422279H1	SATMONN01	g1532072	BLASTN	623	1e-61	88
436	-700455682	700455682H1	SATMON029	g1532047	BLASTN	681	1e-47	77
437	-700477645	700477645H1	SATMON025	g1532048	BLASTX	90	1e-12	71
438	-700550509	700550509H1	SATMON022	g1532047	BLASTN	399	1e-41	80
439	-700572502	700572502H1	SATMON030	g1532048	BLASTX	93	1e-20	70
440	-700618258	700618258H1	SATMON033	g1532072	BLASTN	368	1e-67	92

441	-701165456	701165456H1	SATMONN04	g1532072	BLASTN	436	1e-26	71
442	-L30622289	LIB3062-004-Q1-K1-E10	LIB3062	g1532047	BLASTN	406	1e-26	82
443	-L30623185	LIB3062-026-Q1-K1-E2	LIB3062	g1532072	BLASTN	338	1e-19	62
444	-L30625601	LIB3062-033-Q1-K1-A9	LIB3062	g1403043	BLASTN	758	1e-54	68
445	-L30661842	LIB3066-009-Q1-K1-E4	LIB3066	g1532072	BLASTN	398	1e-22	86
446	-L30681932	LIB3068-020-Q1-K1-C8	LIB3068	g1532072	BLASTN	205	1e-28	89
447	-L30687176	LIB3068-059-Q1-K1-C1	LIB3068	g1532072	BLASTN	216	1e-20	87
448	-L30692075	LIB3069-004-Q1-K1-G11	LIB3069	g1532072	BLASTN	619	1e-40	90
449	-L30783194	LIB3078-052-Q1-K1-E4	LIB3078	g1532072	BLASTN	447	1e-40	76
450	-L30792336	LIB3079-021-Q1-K1-E12	LIB3079	g1532072	BLASTN	217	1e-13	80
451	1471	700106762H1	SATMON010	g1532072	BLASTN	323	1e-59	84
452	1471	701163744H1	SATMONN04	g1532072	BLASTN	220	1e-41	86
453	1471	LIB3059-023-Q1-K1-A11	LIB3059	g1532072	BLASTN	340	1e-39	80
454	1471	700574530H1	SATMON030	g1532072	BLASTN	247	1e-34	82
455	1471	LIB83-003-Q1-E1-G1	LIB83	g1532072	BLASTN	239	1e-24	80
456	1471	701161147H1	SATMONN04	g1532072	BLASTN	239	1e-16	80
457	1471	700477336H1	SATMON025	g1532072	BLASTN	235	1e-10	91
458	16729	700169502H1	SATMON013	g1403043	BLASTN	456	1e-28	65
459	16729	700088201H1	SATMON011	g1532048	BLASTX	123	1e-22	60
460	16729	700165440H1	SATMON013	g1532048	BLASTX	162	1e-15	65
461	16866	700072905H1	SATMON007	g1532047	BLASTN	368	1e-19	71
462	16866	700350558H1	SATMON023	g1532047	BLASTN	301	1e-14	70
463	16866	700088370H1	SATMON011	g1532047	BLASTN	306	1e-14	70
464	2324	LIB3066-042-Q1-K1-H2	LIB3066	g1403043	BLASTN	1344	1e-103	79
465	2324	LIB3059-011-Q1-K1-B2	LIB3059	g1403043	BLASTN	1351	1e-103	78
466	2324	LIB3059-042-Q1-K1-B7	LIB3059	g1403043	BLASTN	1230	1e-100	80
467	2324	LIB3062-036-Q1-K1-F2	LIB3062	g1532072	BLASTN	1097	1e-91	80
468	2324	LIB3069-020-Q1-K1-A5	LIB3069	g1403043	BLASTN	925	1e-82	80
469	2324	LIB189-023-Q1-E1-H3	LIB189	g1532072	BLASTN	1085	1e-81	78
470	2324	700266088H1	SATMON017	g1532047	BLASTN	998	1e-74	80
471	2324	LIB3062-026-Q1-K1-E6	LIB3062	g1532047	BLASTN	982	1e-72	77
472	2324	700083335H1	SATMON011	g1403043	BLASTN	960	1e-71	79
473	2324	LIB189-015-Q1-E1-D3	LIB189	g1532047	BLASTN	970	1e-71	77
474	2324	LIB3079-013-Q1-K1-B3	LIB3079	g1403043	BLASTN	953	1e-70	81
475	2324	700262129H1	SATMON017	g1403043	BLASTN	936	1e-69	79
476	2324	700265667H1	SATMON017	g1403043	BLASTN	694	1e-66	81
477	2324	700807255H1	SATMON036	g1532072	BLASTN	850	1e-62	81
478	2324	700196129H1	SATMON014	g1403043	BLASTN	853	1e-62	83
479	2324	LIB3066-042-Q1-K1-H1	LIB3066	g1403043	BLASTN	842	1e-61	80
480	2324	700458321H1	SATMON029	g1403043	BLASTN	501	1e-58	80
481	2324	700197643H1	SATMON014	g1403043	BLASTN	461	1e-57	80
482	2324	LIB143-029-Q1-E1-H3	LIB143	g1532072	BLASTN	786	1e-56	78
483	2324	700197842H1	SATMON014	g1403043	BLASTN	688	1e-48	85
484	2324	700196807H1	SATMON014	g1532072	BLASTN	678	1e-47	75
485	2324	700263712H1	SATMON017	g1532072	BLASTN	403	1e-45	75
486	2324	LIB143-006-Q1-E1-F9	LIB143	g1403043	BLASTN	324	1e-42	83
487	2324	700267496H1	SATMON017	g1532047	BLASTN	594	1e-40	83
488	2324	700172551H1	SATMON013	g1532072	BLASTN	546	1e-36	74
489	2324	700211893H1	SATMON016	g1532047	BLASTN	542	1e-35	82
490	2324	700264065H1	SATMON017	g1532047	BLASTN	515	1e-34	83
491	2324	700465305H1	SATMON025	g1532073	BLASTX	148	1e-26	61
492	2324	700455052H1	SATMON029	g1403043	BLASTN	274	1e-26	79
493	2324	700473305H1	SATMON025	g1403043	BLASTN	446	1e-26	74
494	2324	700475851H1	SATMON025	g1532047	BLASTN	333	1e-18	78

495	2324	700263111H1	SATMON017	g1532073	BLASTX	165	1e-15	72
496	2324	700465705H1	SATMON025	g1403044	BLASTX	78	1e-8	76
497	3185	700264424H1	SATMON017	g1403043	BLASTN	469	1e-41	81
498	3185	LIB143-007-Q1-E1-H2	LIB143	g1403043	BLASTN	464	1e-36	80
499	3185	700262548H1	SATMON017	g1532047	BLASTN	368	1e-34	80
500	3185	700263845H1	SATMON017	g1403043	BLASTN	455	1e-34	81
501	3185	LIB3068-025-Q1-K1-G10	LIB3068	g1403043	BLASTN	288	1e-32	81
502	3185	700264569H1	SATMON017	g1403043	BLASTN	469	1e-32	82
503	3185	700243571H1	SATMON010	g1532047	BLASTN	461	1e-28	77
504	3185	700265453H1	SATMON017	g1532047	BLASTN	269	1e-27	84
505	3185	700267779H1	SATMON017	g1532047	BLASTN	257	1e-26	85
506	3185	700382373H1	SATMON024	g1532047	BLASTN	255	1e-24	86
507	3185	700258727H1	SATMON017	g1532047	BLASTN	255	1e-24	84
508	3185	LIB3069-018-Q1-K1-F5	LIB3069	g1532047	BLASTN	255	1e-21	77
509	3185	LIB3066-038-Q1-K1-C2	LIB3066	g1403043	BLASTN	242	1e-17	68
510	3185	700334654H1	SATMON019	g1532047	BLASTN	248	1e-17	90
511	3185	700262830H1	SATMON017	g1403043	BLASTN	276	1e-12	78
512	3185	700264727H1	SATMON017	g1532047	BLASTN	255	1e-10	92
513	3185	700238407H1	SATMON010	g1532047	BLASTN	255	1e-10	92
514	3185	700441952H1	SATMON026	g1532047	BLASTN	255	1e-10	92
515	3185	700268188H1	SATMON017	g1532047	BLASTN	255	1e-10	92
516	3185	700801627H1	SATMON036	g1532047	BLASTN	255	1e-10	92
517	3185	700261609H1	SATMON017	g1532047	BLASTN	255	1e-10	92
518	3185	700258510H1	SATMON017	g1532047	BLASTN	255	1e-10	92
519	3185	700258779H1	SATMON017	g1532047	BLASTN	255	1e-10	92
520	3185	700256838H1	SATMON017	g1532047	BLASTN	255	1e-10	92
521	3185	700263361H1	SATMON017	g1532047	BLASTN	255	1e-10	92
522	3185	700257102H1	SATMON017	g1532047	BLASTN	255	1e-10	92
523	3185	700239452H1	SATMON010	g1532047	BLASTN	245	1e-9	91
524	3185	700262045H1	SATMON017	g1532047	BLASTN	250	1e-9	90
525	8	LIB3066-027-Q1-K1-C6	LIB3066	g1532072	BLASTN	1414	1e-181	97
526	8	LIB3066-048-Q1-K1-C9	LIB3066	g1532072	BLASTN	1715	1e-173	98
527	8	LIB3066-007-Q1-K1-B4	LIB3066	g1532072	BLASTN	2146	1e-170	97
528	8	LIB3066-019-Q1-K1-B9	LIB3066	g1532072	BLASTN	1884	1e-168	98
529	8	LIB148-038-Q1-E1-A3	LIB148	g1532072	BLASTN	1585	1e-164	99
530	8	LIB3068-002-Q1-K1-H5	LIB3068	g1532072	BLASTN	2051	1e-162	98
531	8	LIB3067-035-Q1-K1-E9	LIB3067	g1532072	BLASTN	1594	1e-161	98
532	8	LIB189-020-Q1-E1-E8	LIB189	g1532072	BLASTN	1501	1e-160	98
533	8	LIB3078-057-Q1-K1-C12	LIB3078	g1532072	BLASTN	2026	1e-160	96
534	8	LIB3066-053-Q1-K1-A8	LIB3066	g1532072	BLASTN	1713	1e-159	93
535	8	LIB189-006-Q1-E1-C6	LIB189	g1532072	BLASTN	1676	1e-157	98
536	8	LIB3078-054-Q1-K1-E4	LIB3078	g1532072	BLASTN	1997	1e-157	95
537	8	LIB3069-028-Q1-K1-A11	LIB3069	g1532072	BLASTN	1980	1e-156	96
538	8	LIB3069-045-Q1-K1-H7	LIB3069	g1532072	BLASTN	1989	1e-156	98
539	8	LIB148-025-Q1-E1-F7	LIB148	g1532072	BLASTN	1954	1e-154	97
540	8	LIB189-014-Q1-E1-F11	LIB189	g1532072	BLASTN	1925	1e-151	94
541	8	LIB3060-014-Q1-K1-C5	LIB3060	g1532072	BLASTN	1780	1e-150	96
542	8	LIB148-057-Q1-E1-C2	LIB148	g1532072	BLASTN	1000	1e-149	98
543	8	LIB148-015-Q1-E1-F2	LIB148	g1532072	BLASTN	1886	1e-148	94
544	8	LIB3066-045-Q1-K1-A2	LIB3066	g1532072	BLASTN	914	1e-146	91
545	8	LIB148-064-Q1-E1-A3	LIB148	g1532072	BLASTN	1660	1e-146	93
546	8	LIB3061-047-Q1-K1-G1	LIB3061	g1532072	BLASTN	1867	1e-146	92
547	8	LIB148-040-Q1-E1-F1	LIB148	g1532072	BLASTN	1706	1e-145	94
548	8	LIB3069-012-Q1-K1-B6	LIB3069	g1532072	BLASTN	1630	1e-144	87

549	8	LIB3068-009-Q1-K1-A8	LIB3068	g1532072	BLASTN	1802	1e-141	97
550	8	LIB148-004-Q1-E1-D2	LIB148	g1532072	BLASTN	818	1e-139	91
551	8	LIB3068-002-Q1-K1-A1	LIB3068	g1532072	BLASTN	944	1e-137	95
552	8	LIB3067-035-Q1-K1-H9	LIB3067	g1532072	BLASTN	1701	1e-137	98
553	8	LIB3068-033-Q1-K1-G12	LIB3068	g1532072	BLASTN	1093	1e-130	89
554	8	700572229H1	SATMON030	g1532072	BLASTN	1135	1e-128	99
555	8	LIB3078-052-Q1-K1-E1	LIB3078	g1532072	BLASTN	1235	1e-127	85
556	8	700572579H1	SATMON030	g1532072	BLASTN	1625	1e-126	98
557	8	LIB3066-025-Q1-K1-F2	LIB3066	g1532072	BLASTN	1625	1e-126	100
558	8	LIB3068-048-Q1-K1-F9	LIB3068	g1532072	BLASTN	1538	1e-125	98
559	8	700098413H1	SATMON009	g1532072	BLASTN	1595	1e-124	100
560	8	700573235H1	SATMON030	g1532072	BLASTN	1513	1e-123	98
561	8	700090946H1	SATMON011	g1532072	BLASTN	1585	1e-123	100
562	8	700092465H1	SATMON008	g1532072	BLASTN	1541	1e-122	98
563	8	700074625H1	SATMON007	g1532072	BLASTN	1570	1e-122	100
564	8	LIB3059-007-Q1-K1-C10	LIB3059	g1532072	BLASTN	1401	1e-119	94
565	8	700072828H1	SATMON007	g1532072	BLASTN	1540	1e-119	100
566	8	700619106H1	SATMON034	g1532072	BLASTN	833	1e-118	97
567	8	700074853H1	SATMON007	g1532072	BLASTN	1515	1e-117	100
568	8	700201293H1	SATMON003	g1532072	BLASTN	1517	1e-117	97
569	8	700075896H1	SATMON007	g1532072	BLASTN	1006	1e-115	99
570	8	LIB3059-052-Q1-K1-A1	LIB3059	g1532072	BLASTN	1241	1e-115	92
571	8	700091576H1	SATMON011	g1532072	BLASTN	943	1e-114	98
572	8	LIB3078-015-Q1-K1-D7	LIB3078	g1532072	BLASTN	1218	1e-114	87
573	8	700074733H1	SATMON007	g1532072	BLASTN	1475	1e-114	100
574	8	700381421H1	SATMON023	g1532072	BLASTN	1475	1e-114	100
575	8	700085594H1	SATMON011	g1532072	BLASTN	1477	1e-114	99
576	8	700095883H1	SATMON008	g1532072	BLASTN	1477	1e-114	99
577	8	700338237H1	SATMON020	g1532072	BLASTN	1478	1e-114	99
578	8	700549813H1	SATMON022	g1532072	BLASTN	1481	1e-114	99
579	8	700097935H1	SATMON009	g1532072	BLASTN	1484	1e-114	98
580	8	700572978H1	SATMON030	g1532072	BLASTN	865	1e-113	96
581	8	700196464H1	SATMON014	g1532072	BLASTN	1044	1e-113	92
582	8	700381412H1	SATMON023	g1532072	BLASTN	1168	1e-113	98
583	8	700027839H1	SATMON003	g1532072	BLASTN	1472	1e-113	99
584	8	700623344H1	SATMON034	g1532072	BLASTN	1085	1e-112	94
585	8	700025858H1	SATMON003	g1532072	BLASTN	1455	1e-112	100
586	8	LIB3059-017-Q1-K1-H5	LIB3059	g1532072	BLASTN	1459	1e-112	97
587	8	700475019H1	SATMON025	g1532072	BLASTN	1390	1e-111	100
588	8	700338357H1	SATMON020	g1532072	BLASTN	1440	1e-111	100
589	8	700256796H1	SATMON017	g1532072	BLASTN	1400	1e-110	100
590	8	700071692H1	SATMON007	g1532072	BLASTN	1430	1e-110	98
591	8	700339073H1	SATMON020	g1532072	BLASTN	1357	1e-109	95
592	8	700106916H1	SATMON010	g1532072	BLASTN	1415	1e-109	93
593	8	700468234H1	SATMON025	g1532072	BLASTN	1420	1e-109	100
594	8	700214482H1	SATMON016	g1532072	BLASTN	1425	1e-109	100
595	8	700466437H1	SATMON025	g1532072	BLASTN	1403	1e-108	98
596	8	700205576H1	SATMON003	g1532072	BLASTN	1404	1e-108	98
597	8	700043455H1	SATMON004	g1532072	BLASTN	1405	1e-108	100
598	8	700348439H1	SATMON023	g1532072	BLASTN	1407	1e-108	99
599	8	700093131H1	SATMON008	g1532072	BLASTN	755	1e-107	100
600	8	700571851H1	SATMON030	g1532072	BLASTN	1302	1e-107	99
601	8	700088142H1	SATMON011	g1532072	BLASTN	1392	1e-107	99
602	8	700085934H1	SATMON011	g1532072	BLASTN	1397	1e-107	98

603	8	700028465H1	SATMON003	g1532072	BLASTN	1258	1e-106	98
604	8	700236818H1	SATMON010	g1532072	BLASTN	1380	1e-106	100
605	8	700583691H1	SATMON031	g1532072	BLASTN	1382	1e-106	99
606	8	700095585H1	SATMON008	g1532072	BLASTN	1383	1e-106	94
607	8	700105140H1	SATMON010	g1532072	BLASTN	1375	1e-105	100
608	8	700338486H1	SATMON020	g1532072	BLASTN	893	1e-104	98
609	8	700214178H1	SATMON016	g1532072	BLASTN	1361	1e-104	99
610	8	700090613H1	SATMON011	g1532072	BLASTN	699	1e-103	99
611	8	700576189H1	SATMON030	g1532072	BLASTN	1254	1e-103	93
612	8	700475734H1	SATMON025	g1532072	BLASTN	1256	1e-103	99
613	8	700028218H1	SATMON003	g1532072	BLASTN	1275	1e-103	100
614	8	700088644H1	SATMON011	g1532072	BLASTN	1351	1e-103	98
615	8	700378768H1	SATMON020	g1532072	BLASTN	1067	1e-102	98
616	8	LIB3067-035-Q1-K1-H10	LIB3067	g1532072	BLASTN	1233	1e-102	95
617	8	700043466H1	SATMON004	g1532072	BLASTN	1331	1e-102	97
618	8	LIB148-057-Q1-E1-C3	LIB148	g1532072	BLASTN	1283	1e-101	92
619	8	700217574H1	SATMON016	g1532072	BLASTN	1320	1e-101	100
620	8	700042332H1	SATMON004	g1532072	BLASTN	1321	1e-101	97
621	8	700440926H1	SATMON026	g1532072	BLASTN	1323	1e-101	99
622	8	700552284H1	SATMON022	g1532072	BLASTN	1329	1e-101	97
623	8	700216613H1	SATMON016	g1532072	BLASTN	689	1e-100	97
624	8	LIB3067-054-Q1-K1-F6	LIB3067	g1532072	BLASTN	1030	1e-100	94
625	8	LIB3060-038-Q1-K1-B9	LIB3060	g1532072	BLASTN	1108	1e-100	92
626	8	700218755H1	SATMON011	g1532072	BLASTN	1116	1e-100	99
627	8	700338042H1	SATMON020	g1532072	BLASTN	1270	1e-100	99
628	8	700268009H1	SATMON017	g1532072	BLASTN	1307	1e-100	92
629	8	700578491H1	SATMON031	g1532072	BLASTN	1309	1e-100	97
630	8	700030037H1	SATMON003	g1532072	BLASTN	1310	1e-100	97
631	8	700221406H1	SATMON011	g1532072	BLASTN	1315	1e-100	100
632	8	700157357H1	SATMON012	g1532072	BLASTN	1315	1e-100	98
633	8	700476926H1	SATMON025	g1532072	BLASTN	680	1e-99	97
634	8	700475068H1	SATMON025	g1532072	BLASTN	792	1e-99	98
635	8	700469741H1	SATMON025	g1532072	BLASTN	831	1e-99	99
636	8	700082377H1	SATMON011	g1532072	BLASTN	1081	1e-99	99
637	8	700217143H1	SATMON016	g1532072	BLASTN	1208	1e-99	98
638	8	700339433H1	SATMON020	g1532072	BLASTN	1295	1e-99	100
639	8	700196627H1	SATMON014	g1532072	BLASTN	1295	1e-99	100
640	8	700259354H1	SATMON017	g1532072	BLASTN	1305	1e-99	92
641	8	700610842H1	SATMON022	g1532072	BLASTN	843	1e-98	97
642	8	700218059H1	SATMON016	g1532072	BLASTN	1036	1e-98	99
643	8	700421727H1	SATMONN01	g1532072	BLASTN	1172	1e-98	97
644	8	700158660H1	SATMON012	g1532072	BLASTN	1285	1e-98	100
645	8	700806856H1	SATMON036	g1532072	BLASTN	1194	1e-97	97
646	8	700583512H1	SATMON031	g1532072	BLASTN	1219	1e-97	97
647	8	700025502H1	SATMON004	g1532072	BLASTN	1276	1e-97	99
648	8	700159562H1	SATMON012	g1532072	BLASTN	1277	1e-97	99
649	8	700223731H1	SATMON011	g1532072	BLASTN	1277	1e-97	99
650	8	700156494H1	SATMON012	g1532072	BLASTN	1280	1e-97	100
651	8	700160533H1	SATMON012	g1532072	BLASTN	1281	1e-97	97
652	8	LIB3066-055-Q1-K1-D8	LIB3066	g1532072	BLASTN	672	1e-96	99
653	8	700405152H1	SATMON028	g1532072	BLASTN	828	1e-96	98
654	8	LIB148-040-Q1-E1-A8	LIB148	g1532072	BLASTN	1268	1e-96	83
655	8	700438437H1	SATMON026	g1532072	BLASTN	1246	1e-95	99
656	8	700267245H1	SATMON017	g1532072	BLASTN	1248	1e-95	93



657	8	700156129H2	SATMON007	g1532072	BLASTN	1250	1e-95	100
658	8	700203246H1	SATMON003	g1532072	BLASTN	1250	1e-95	100
659	8	700168339H1	SATMON013	g1532072	BLASTN	915	1e-94	98
660	8	LIB3067-036-Q1-K1-F9	LIB3067	g1532072	BLASTN	1110	1e-94	95
661	8	700193769H1	SATMON014	g1532072	BLASTN	1240	1e-94	100
662	8	700094014H1	SATMON008	g1532072	BLASTN	1241	1e-94	91
663	8	700020311H1	SATMON001	g1532072	BLASTN	1228	1e-93	99
664	8	700195083H1	SATMON014	g1532072	BLASTN	1229	1e-93	98
665	8	700193901H1	SATMON014	g1532072	BLASTN	1229	1e-93	98
666	8	700194639H1	SATMON014	g1532072	BLASTN	1231	1e-93	99
667	8	700350117H1	SATMON023	g1532072	BLASTN	1143	1e-92	99
668	8	700239867H1	SATMON010	g1532072	BLASTN	1210	1e-92	98
669	8	LIB3069-028-Q1-K1-D1	LIB3069	g1532072	BLASTN	1217	1e-92	94
670	8	700355756H1	SATMON024	g1532072	BLASTN	618	1e-91	96
671	8	700265492H1	SATMON017	g1532072	BLASTN	620	1e-91	94
672	8	700579021H1	SATMON031	g1532072	BLASTN	656	1e-91	96
673	8	LIB3079-015-Q1-K1-B11	LIB3079	g1532072	BLASTN	1008	1e-91	83
674	8	700572888H2	SATMON030	g1532072	BLASTN	1159	1e-91	99
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676	8	700085757H1	SATMON011	g1532072	BLASTN	1200	1e-91	100
677	8	700162756H1	SATMON013	g1532072	BLASTN	1208	1e-91	99
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679	8	700224378H1	SATMON011	g1532072	BLASTN	696	1e-90	99
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681	8	700159338H1	SATMON012	g1532072	BLASTN	1193	1e-90	97
682	8	700169217H1	SATMON013	g1532072	BLASTN	1196	1e-90	99
683	8	700551548H1	SATMON022	g1532072	BLASTN	839	1e-89	96
684	8	700197059H1	SATMON014	g1532072	BLASTN	1025	1e-89	95
685	8	700469834H1	SATMON025	g1532072	BLASTN	650	1e-88	97
686	8	700569778H1	SATMON030	g1532072	BLASTN	682	1e-88	91
687	8	700194845H1	SATMON014	g1532072	BLASTN	1171	1e-88	99
688	8	700445861H1	SATMON027	g1532072	BLASTN	536	1e-87	99
689	8	700100536H1	SATMON009	g1532072	BLASTN	666	1e-87	94
690	8	701163801H1	SATMONN04	g1532072	BLASTN	805	1e-87	94
691	8	LIB3060-035-Q1-K1-E3	LIB3060	g1532072	BLASTN	922	1e-87	95
692	8	700170872H1	SATMON013	g1532072	BLASTN	943	1e-87	97
693	8	700241270H1	SATMON010	g1532072	BLASTN	1066	1e-86	96
694	8	700457981H1	SATMON029	g1532072	BLASTN	1140	1e-86	92
695	8	700158339H1	SATMON012	g1532072	BLASTN	1142	1e-86	99
696	8	700019442H1	SATMON001	g1532072	BLASTN	1145	1e-86	100
697	8	700149885H1	SATMON007	g1532072	BLASTN	1133	1e-85	99
698	8	700018255H1	SATMON001	g1532072	BLASTN	1135	1e-85	100
699	8	700244026H1	SATMON010	g1532072	BLASTN	1044	1e-84	89
700	8	LIB3060-027-Q1-K1-B2	LIB3060	g1532072	BLASTN	1117	1e-84	99
701	8	700170960H1	SATMON013	g1532072	BLASTN	696	1e-83	99
702	8	700267487H1	SATMON017	g1532072	BLASTN	1061	1e-83	92
703	8	700152754H1	SATMON007	g1532072	BLASTN	1107	1e-83	99
704	8	700378260H1	SATMON019	g1532072	BLASTN	1112	1e-83	92
705	8	700455089H1	SATMON029	g1532072	BLASTN	583	1e-82	97
706	8	700167322H1	SATMON013	g1532072	BLASTN	1080	1e-81	98
707	8	LIB3060-035-Q1-K1-H1	LIB3060	g1532072	BLASTN	782	1e-80	94
708	8	700204067H1	SATMON003	g1532072	BLASTN	1011	1e-80	98
709	8	700049271H1	SATMON003	g1532072	BLASTN	627	1e-79	93
710	8	700442065H1	SATMON026	g1532072	BLASTN	1043	1e-78	91



765	8	700431071H1	SATMONN01	g1532072	BLASTN	366	1e-50	94
766	8	700171582H1	SATMON013	g1532072	BLASTN	492	1e-50	99
767	8	700450579H1	SATMON028	g1532072	BLASTN	338	1e-49	97
768	8	700084149H1	SATMON011	g1532072	BLASTN	672	1e-47	98
769	8	700095070H1	SATMON008	g1532072	BLASTN	662	1e-46	98
770	8	700570827H1	SATMON030	g1532072	BLASTN	344	1e-44	83
771	8	LIB3061-057-Q1-K1-G12	LIB3061	g1532072	BLASTN	378	1e-43	75
772	8	700194918H1	SATMON014	g1532072	BLASTN	626	1e-43	89
773	8	700378894H1	SATMON020	g1532072	BLASTN	495	1e-42	97
774	8	700468029H1	SATMON025	g1532072	BLASTN	616	1e-42	98
775	8	LIB143-012-Q1-E1-H8	LIB143	g1532072	BLASTN	637	1e-42	98
776	8	LIB3060-037-Q1-K1-H4	LIB3060	g1532072	BLASTN	638	1e-42	86
777	8	700433327H1	SATMONN01	g1532072	BLASTN	365	1e-41	86
778	8	700623611H1	SATMON034	g1532072	BLASTN	316	1e-39	95
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780	8	700616273H1	SATMON033	g1532072	BLASTN	534	1e-38	98
781	8	700464702H1	SATMON025	g1532072	BLASTN	566	1e-38	99
782	8	700338809H1	SATMON020	g1532072	BLASTN	550	1e-37	100
783	8	700092622H1	SATMON008	g1532072	BLASTN	561	1e-37	99
784	8	LIB3060-043-Q1-K1-A10	LIB3060	g1532072	BLASTN	352	1e-36	93
785	8	700265611H1	SATMON017	g1532072	BLASTN	548	1e-36	95
786	8	700100319H1	SATMON009	g1532072	BLASTN	552	1e-36	98
787	8	700092696H1	SATMON008	g1532072	BLASTN	552	1e-36	98
788	8	700076750H1	SATMON007	g1532072	BLASTN	526	1e-35	99
789	8	700082896H1	SATMON011	g1532072	BLASTN	526	1e-35	99
790	8	700075411H1	SATMON007	g1532072	BLASTN	319	1e-34	91
791	8	701178051H1	SATMONN05	g1532072	BLASTN	342	1e-34	94
792	8	700266358H1	SATMON017	g1532072	BLASTN	520	1e-34	95
793	8	700453981H1	SATMON029	g1532072	BLASTN	505	1e-33	96
794	8	700584238H1	SATMON031	g1532072	BLASTN	509	1e-33	91
795	8	700103871H1	SATMON010	g1532072	BLASTN	491	1e-32	99
796	8	700264460H1	SATMON017	g1532072	BLASTN	495	1e-32	95
797	8	700205122H1	SATMON003	g1532072	BLASTN	501	1e-32	99
798	8	700165768H1	SATMON013	g1532072	BLASTN	420	1e-31	98
799	8	700077179H1	SATMON007	g1532072	BLASTN	471	1e-30	98
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803	8	700088193H1	SATMON011	g1532072	BLASTN	488	1e-30	97
804	8	700214511H1	SATMON016	g1532072	BLASTN	298	1e-29	93
805	8	700257186H1	SATMON017	g1532072	BLASTN	465	1e-29	95
806	8	700335814H1	SATMON019	g1532072	BLASTN	469	1e-29	93
807	8	700236166H1	SATMON010	g1532072	BLASTN	450	1e-28	95
808	8	700468243H1	SATMON025	g1532072	BLASTN	456	1e-28	98
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812	8	700264846H1	SATMON017	g1532072	BLASTN	430	1e-25	94
813	8	700266803H1	SATMON017	g1532072	BLASTN	293	1e-24	75
814	8	700086134H1	SATMON011	g1532072	BLASTN	422	1e-24	97
815	8	700162001H1	SATMON012	g1532072	BLASTN	279	1e-23	96
816	8	700044825H1	SATMON004	g1532072	BLASTN	411	1e-23	98
817	8	700074679H1	SATMON007	g1532072	BLASTN	411	1e-23	98
818	8	700267694H1	SATMON017	g1532072	BLASTN	320	1e-22	92





2549	16	700872967H1	SOYMON018	g1421750	BLASTN	704	1e-49	75
2550	16	701119960H1	SOYMON037	g1421750	BLASTN	684	1e-48	81
2551	16	700873306H1	SOYMON018	g1421750	BLASTN	693	1e-48	83
2552	16	700751484H1	SOYMON014	g1421750	BLASTN	635	1e-47	82
2553	16	700958865H1	SOYMON022	g1421750	BLASTN	486	1e-46	82
2554	16	700872833H1	SOYMON018	g1421750	BLASTN	659	1e-46	84
2555	16	700871673H1	SOYMON018	g1421750	BLASTN	659	1e-46	84
2556	16	700872801H1	SOYMON018	g1421750	BLASTN	659	1e-46	84
2557	16	700847572H1	SOYMON021	g1421750	BLASTN	660	1e-46	75
2558	16	700728046H1	SOYMON009	g1421750	BLASTN	661	1e-46	84
2559	16	700894721H1	SOYMON024	g1421750	BLASTN	550	1e-43	84
2560	16	700730946H1	SOYMON009	g1421750	BLASTN	307	1e-42	77
2561	16	700727943H1	SOYMON009	g1421750	BLASTN	378	1e-42	79
2562	16	701101238H1	SOYMON028	g1421750	BLASTN	456	1e-42	92
2563	16	700758519H1	SOYMON015	g1421750	BLASTN	358	1e-41	77
2564	16	700740343H1	SOYMON012	g1531764	BLASTN	607	1e-41	72
2565	16	700848076H1	SOYMON021	g1531764	BLASTN	469	1e-40	89
2566	16	LIB3051-080-Q1-K1-H7	LIB3051	g1531764	BLASTN	472	1e-38	90
2567	16	700955916H1	SOYMON022	g1531764	BLASTN	427	1e-37	90
2568	16	700873876H1	SOYMON018	g1421750	BLASTN	465	1e-37	80
2569	16	700746271H1	SOYMON013	g1421750	BLASTN	515	1e-37	80
2570	16	700662887H1	SOYMON005	g1421750	BLASTN	506	1e-35	82
2571	16	700983591H1	SOYMON009	g1421750	BLASTN	537	1e-35	82
2572	16	700752343H1	SOYMON014	g1421750	BLASTN	488	1e-34	73
2573	16	700901973H1	SOYMON027	g1421750	BLASTN	490	1e-34	82
2574	16	LIB3053-013-Q1-N1-B11	LIB3053	g1421750	BLASTN	522	1e-34	75
2575	16	700988481H1	SOYMON009	g1421750	BLASTN	506	1e-33	78
2576	16	LIB3051-078-Q1-K1-C12	LIB3051	g1421750	BLASTN	302	1e-32	75
2577	16	700985640H1	SOYMON009	g1421750	BLASTN	449	1e-30	86
2578	16	701127038H1	SOYMON037	g1421750	BLASTN	467	1e-30	67
2579	16	700898677H1	SOYMON027	g1531764	BLASTN	474	1e-30	89
2580	16	700742260H1	SOYMON012	g1421750	BLASTN	476	1e-30	70
2581	16	701103203H1	SOYMON028	g1421750	BLASTN	287	1e-27	81
2582	16	700897144H1	SOYMON027	g1421750	BLASTN	264	1e-26	75
2583	16	700832039H1	SOYMON019	g1421750	BLASTN	390	1e-26	82
2584	16	700743311H1	SOYMON012	g1421750	BLASTN	429	1e-26	87
2585	16	LIB3040-044-Q1-E1-B8	LIB3040	g1421750	BLASTN	282	1e-25	92
2586	16	LIB3051-024-Q1-K1-C4	LIB3051	g2394382	BLASTX	110	1e-24	95
2587	16	701011765H1	SOYMON019	g1421750	BLASTN	383	1e-21	76
2588	16	700893428H1	SOYMON024	g1421752	BLASTX	106	1e-20	82
2589	16	700996204H1	SOYMON018	g1421750	BLASTN	288	1e-20	75
2590	16	LIB3049-028-Q1-E1-C6	LIB3049	g1421750	BLASTN	230	1e-19	90
2591	16	701103628H1	SOYMON028	g1421750	BLASTN	277	1e-18	73
2592	16	700972636H1	SOYMON005	g1421750	BLASTN	286	1e-16	73
2593	16	700875417H1	SOYMON018	g1421750	BLASTN	209	1e-15	75
2594	16	701207702H1	SOYMON035	g1421752	BLASTX	153	1e-14	83
2595	16	701054556H1	SOYMON032	g1421750	BLASTN	230	1e-14	75
2596	16	701046407H1	SOYMON032	g1421750	BLASTN	230	1e-14	75
2597	16	701117634H1	SOYMON037	g1421750	BLASTN	230	1e-14	74
2598	16	701127714H1	SOYMON037	g1421750	BLASTN	230	1e-14	75
2599	16	700663239H1	SOYMON005	g1421750	BLASTN	210	1e-13	74
2600	16	700738510H1	SOYMON012	g1421750	BLASTN	212	1e-13	72
2601	16	700943559H1	SOYMON024	g1421750	BLASTN	215	1e-13	71
2602	16	700666278H1	SOYMON005	g1421750	BLASTN	290	1e-13	93

2603	16	700663144H1	SOYMON005	g1421750	BLASTN	204	1e-12	74
2604	16	700684147H1	SOYMON008	g1421750	BLASTN	278	1e-12	89
2605	16	700953076H1	SOYMON022	g1421750	BLASTN	285	1e-12	92
2606	16	701099931H1	SOYMON028	g1421750	BLASTN	202	1e-11	72
2607	16	701058910H1	SOYMON033	g1421750	BLASTN	216	1e-11	74
2608	16	700754949H1	SOYMON014	g1421750	BLASTN	266	1e-11	89
2609	16	700562967H1	SOYMON002	g1421750	BLASTN	266	1e-11	82
2610	16	701012790H1	SOYMON019	g1421750	BLASTN	251	1e-10	89
2611	16	701101739H1	SOYMON028	g1421750	BLASTN	260	1e-10	82
2612	16	701118211H1	SOYMON037	g1490554	BLASTX	97	1e-9	47
2613	16	700844394H1	SOYMON021	g1917013	BLASTX	116	1e-9	85
2614	16	701048560H1	SOYMON032	g1421750	BLASTN	198	1e-9	79
2615	16	701062373H1	SOYMON033	g1421750	BLASTN	203	1e-9	71
2616	16	701120496H1	SOYMON037	g1421750	BLASTN	230	1e-9	82
2617	16	701049680H1	SOYMON032	g1421750	BLASTN	177	1e-8	70
2618	16	700748760H1	SOYMON013	g1421750	BLASTN	230	1e-8	91
2619	16	700889482H1	SOYMON024	g1421750	BLASTN	235	1e-8	91
2620	16	700725013H1	SOYMON009	g1421750	BLASTN	239	1e-8	85
2621	16048	LIB3028-004-Q1-B1-E2	LIB3028	g1421750	BLASTN	879	1e-64	68
2622	16048	700761667H1	SOYMON015	g1421750	BLASTN	528	1e-35	71
2623	16048	700958003H1	SOYMON022	g1421750	BLASTN	428	1e-26	78

#### ASPARTATE KINASE ( EC 2.7.2.4 )

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
858	-700018870	700018870H1	SATMON001	g500850	BLASTN	1095	1e-82	100
859	-700085903	700085903H1	SATMON011	g500852	BLASTN	1606	1e-124	99
860	-700086169	700086169H1	SATMON011	g500852	BLASTN	559	1e-94	92
861	-700096679	700096679H1	SATMON008	g500850	BLASTN	1131	1e-85	99
862	-700096794	700096794H1	SATMON008	g500850	BLASTN	1515	1e-117	100
863	-700106390	700106390H1	SATMON010	g2243115	BLASTN	544	1e-51	72
864	-700168286	700168286H1	SATMON013	g2243115	BLASTN	519	1e-34	72
865	-700171363	700171363H1	SATMON013	g500850	BLASTN	1085	1e-81	94
866	-700194781	700194781H1	SATMON014	g2257742	BLASTN	635	1e-44	79
867	-700213839	700213839H1	SATMON016	g500850	BLASTN	664	1e-75	94
868	-700219756	700219756H1	SATMON011	g500850	BLASTN	1359	1e-104	99
869	-700258808	700258808H1	SATMON017	g500850	BLASTN	630	1e-85	96
870	-700263439	700263439H1	SATMON017	g2243115	BLASTN	448	1e-43	76
871	-700266615	700266615H1	SATMON017	g2243116	BLASTX	179	1e-17	73
872	-700342655	700342655H1	SATMON021	g2257743	BLASTX	213	1e-22	82
873	-700343285	700343285H1	SATMON021	g2257742	BLASTN	437	1e-25	66
874	-700467533	700467533H1	SATMON025	g2243115	BLASTN	439	1e-26	79
875	-700548678	700548678H1	SATMON022	g147979	BLASTX	147	1e-13	62
876	-700613618	700613618H1	SATMON033	g500850	BLASTN	965	1e-81	98
877	-L30691987	LIB3069-018-Q1-K1-A3	LIB3069	g2243115	BLASTN	266	1e-10	57
878	12201	700457103H1	SATMON029	g2257742	BLASTN	789	1e-56	76
879	12201	700457111H1	SATMON029	g2243115	BLASTN	513	1e-33	76
880	12931	700380864H1	SATMON023	g500852	BLASTN	1450	1e-111	95
881	12931	700105610H1	SATMON010	g500852	BLASTN	1046	1e-105	95
882	12931	700380848H1	SATMON023	g500852	BLASTN	1193	1e-96	95
883	12931	700205392H1	SATMON003	g500852	BLASTN	1065	1e-94	93
884	12931	700552314H1	SATMON022	g500852	BLASTN	908	1e-86	90
885	12931	700551915H1	SATMON022	g500852	BLASTN	1090	1e-81	90





2654 30187 700556105H1 SOYMON001 g1359593 BLASTX 132 1e-11 56

**O-SUCCINYLMOMOSERINE (THIOL)-LYASE ( EC 4.2.99.9 )**

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
905	-700049526	700049526H1	SATMON003	g2198852	BLASTN	251	1e-9	76
906	-700086788	700086788H1	SATMON011	g2198850	BLASTN	631	1e-61	88
907	-700460589	700460589H1	SATMON030	g2198850	BLASTN	251	1e-32	83
908	-700577561	700577561H1	SATMON031	g2198852	BLASTN	207	1e-15	82
909	-700579840	700579840H1	SATMON031	g2198850	BLASTN	216	1e-12	94
910	-700616013	700616013H1	SATMON033	g2198852	BLASTN	278	1e-39	81
911	-L1892785	LIB189-011-Q1-E1-A10	LIB189	g2198852	BLASTN	311	1e-14	82
912	-L30594402	LIB3059-042-Q1-K1-E11	LIB3059	g2198852	BLASTN	489	1e-40	84
913	-L30604293	LIB3060-028-Q1-K1-E7	LIB3060	g2198852	BLASTN	380	1e-20	83
914	-L30693289	LIB3069-016-Q1-K1-G10	LIB3069	g2198852	BLASTN	197	1e-10	79
915	10571	700224881H1	SATMON011	g2198852	BLASTN	319	1e-15	75
916	10801	700429049H1	SATMONN01	g2198852	BLASTN	210	1e-16	82
917	10801	700167813H1	SATMON013	g2198852	BLASTN	200	1e-15	82
918	10801	700074027H1	SATMON007	g2198852	BLASTN	178	1e-11	80
919	16379	LIB3061-035-Q1-K1-B6	LIB3061	g2198850	BLASTN	2184	1e-173	99
920	16379	700259309H1	SATMON017	g2198850	BLASTN	1259	1e-108	92
921	16379	700051696H1	SATMON003	g2198850	BLASTN	1392	1e-107	96
922	16379	700239553H1	SATMON010	g2198850	BLASTN	1265	1e-96	96
923	16379	700042846H1	SATMON004	g2198850	BLASTN	1208	1e-91	95
924	16379	700092741H1	SATMON008	g2198850	BLASTN	1042	1e-89	95
925	16379	LIB3061-017-Q1-K1-D7	LIB3061	g2198850	BLASTN	1183	1e-89	99
926	16379	700206765H1	SATMON003	g2198850	BLASTN	1095	1e-82	81
927	16379	700150281H1	SATMON007	g2198850	BLASTN	1024	1e-76	94
928	16379	700165862H1	SATMON013	g2198850	BLASTN	948	1e-75	94
929	2221	LIB3060-017-Q1-K1-B10	LIB3060	g2198850	BLASTN	1819	1e-161	99
930	2221	LIB84-006-Q1-E1-F3	LIB84	g2198852	BLASTN	1615	1e-153	97
931	2221	700575334H1	SATMON030	g2198850	BLASTN	1570	1e-124	99
932	2221	700206250H1	SATMON003	g2198850	BLASTN	1515	1e-117	100
933	2221	700095073H1	SATMON008	g2198852	BLASTN	1490	1e-115	100
934	2221	700571230H1	SATMON030	g2198850	BLASTN	1355	1e-114	93
935	2221	700157358H1	SATMON012	g2198850	BLASTN	1370	1e-105	100
936	2221	700379811H1	SATMON021	g2198850	BLASTN	1345	1e-103	93
937	2221	700041570H1	SATMON004	g2198850	BLASTN	1325	1e-101	100
938	2221	700104063H1	SATMON010	g2198850	BLASTN	1157	1e-95	90
939	2221	700378265H1	SATMON019	g2198850	BLASTN	771	1e-94	99
940	2221	700235329H1	SATMON010	g2198850	BLASTN	1234	1e-94	93
941	2221	LIB3068-057-Q1-K1-D1	LIB3068	g2198852	BLASTN	1209	1e-91	94
942	2221	700159158H1	SATMON012	g2198850	BLASTN	1174	1e-89	94
943	2221	700580854H1	SATMON031	g2198850	BLASTN	754	1e-86	92
944	2221	700623409H1	SATMON034	g2198850	BLASTN	910	1e-84	96
945	2221	701164706H1	SATMONN04	g2198852	BLASTN	577	1e-83	94
946	2221	700164719H1	SATMON013	g2198852	BLASTN	831	1e-83	99
947	2221	700158146H1	SATMON012	g2198850	BLASTN	1100	1e-82	93
948	2221	700203970H1	SATMON003	g2198852	BLASTN	996	1e-80	99
949	2221	700158313H1	SATMON012	g2198850	BLASTN	1036	1e-77	93
950	2221	700425211H1	SATMONN01	g2198852	BLASTN	485	1e-61	96
951	2221	700167764H1	SATMON013	g2198850	BLASTN	793	1e-57	93
952	23788	700102780H1	SATMON010	g2198852	BLASTN	1131	1e-104	99

953	23788	701167458H1	SATMONN05	g2198852	BLASTN	1069	1e-90	93
2655	-700900206	700900206H1	SOYMON027	g1742961	BLASTX	215	1e-24	78
2656	-GM40351	LIB3051-114-Q1-K1-H12	LIB3051	g2198851	BLASTX	122	1e-25	96
2657	12502	701101592H1	SOYMON028	g146846	BLASTX	103	1e-18	44
2658	12502	701106834H1	SOYMON036	g146846	BLASTX	103	1e-18	44
2659	13820	LIB3055-003-Q1-N1-F12	LIB3055	g3202028	BLASTX	193	1e-35	94
2660	8119	700989656H1	SOYMON011	g1742960	BLASTN	815	1e-59	79

#### CYSTATHIONINE $\beta$ -LYASE ( EC 4.4.1.8 )

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
954	-700155172	700155172H1	SATMON007	g704397	BLASTX	361	1e-43	78
955	-L362943	LIB36-013-Q1-E1-G10	LIB36	g704396	BLASTN	818	1e-59	75
956	19856	700240664H1	SATMON010	g704396	BLASTN	496	1e-31	69
957	19856	700572496H1	SATMON030	g704396	BLASTN	446	1e-26	68
958	22960	701170964H1	SATMONN05	g704396	BLASTN	778	1e-56	78
959	22960	LIB3061-002-Q1-K2-F9	LIB3061	g704396	BLASTN	765	1e-53	77
960	22960	701172780H2	SATMONN05	g704396	BLASTN	686	1e-48	73
961	22960	700578571H1	SATMON031	g704397	BLASTX	222	1e-23	89
962	30752	LIB3078-055-Q1-K1-C8	LIB3078	g704396	BLASTN	747	1e-51	71
963	30752	700086603H1	SATMON011	g704397	BLASTX	192	1e-18	64
2661	-701001147	701001147H1	SOYMON018	g704396	BLASTN	847	1e-61	78
2662	18602	700566066H1	SOYMON002	g704396	BLASTN	751	1e-57	79
2663	18602	700890955H1	SOYMON024	g704396	BLASTN	698	1e-49	77
2664	18602	700896865H1	SOYMON027	g704396	BLASTN	682	1e-48	77
2665	5144	LIB3050-006-Q1-E1-A9	LIB3050	g1399263	BLASTX	96	1e-31	41

#### 5-METHYLTETRAHYDROPTEROYLTRIGLUTAMATE--HOMOCYSTEINE S-METHYLTRANSFERASE ( EC 2.1.1.14 )

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
964	-700212217	700212217H1	SATMON016	g886471	BLASTX	151	1e-16	93
965	-700333966	700333966H1	SATMON019	g886470	BLASTN	370	1e-23	66
966	-700377403	700377403H1	SATMON019	g2738247	BLASTN	305	1e-26	80
967	-700571893	700571893H1	SATMON030	g974781	BLASTN	262	1e-20	80
968	-701165656	701165656H1	SATMONN04	g2738247	BLASTN	450	1e-50	73
969	-L30622954	LIB3062-030-Q1-K1-C9	LIB3062	g886470	BLASTN	366	1e-26	81
970	-L30662004	LIB3066-026-Q1-K1-F1	LIB3066	g2738248	BLASTX	140	1e-28	77
971	-L30671450	LIB3067-001-Q1-K1-C6	LIB3067	g2738247	BLASTN	553	1e-46	69
972	-L30694410	LIB3069-057-Q1-K1-B4	LIB3069	g886470	BLASTN	447	1e-26	67
973	1	700452404H1	SATMON028	g453939	BLASTX	59	1e-17	94
974	13513	700049012H1	SATMON003	g1814402	BLASTN	1006	1e-74	81
975	13513	700086058H1	SATMON011	g1814402	BLASTN	982	1e-72	81
976	13513	700235814H1	SATMON010	g1814402	BLASTN	890	1e-65	79
977	13513	700170015H1	SATMON013	g1814402	BLASTN	578	1e-39	78
978	3835	700093161H1	SATMON008	g974781	BLASTN	858	1e-62	76
979	3835	700223321H1	SATMON011	g974781	BLASTN	836	1e-60	80
980	3835	700454003H1	SATMON029	g974781	BLASTN	823	1e-59	82
981	3835	700238925H1	SATMON010	g2738247	BLASTN	733	1e-55	74
982	3835	700075142H1	SATMON007	g2738247	BLASTN	569	1e-52	73
983	3835	700151969H1	SATMON007	g974781	BLASTN	660	1e-46	76
984	3835	700281434H2	SATMON019	g974781	BLASTN	560	1e-45	79

985	3835	700084914H1	SATMON011	g974781	BLASTN	440	1e-36	79
986	3835	700215135H1	SATMON016	g974781	BLASTN	543	1e-36	78
987	3835	700202218H1	SATMON003	g974781	BLASTN	389	1e-23	78
988	3835	700281467H2	SATMON019	g2738248	BLASTX	123	1e-12	71
989	456	LIB3059-019-Q1-K1-G6	LIB3059	g974781	BLASTN	1493	1e-115	83
990	456	LIB3068-012-Q1-K1-G7	LIB3068	g974781	BLASTN	1471	1e-113	81
991	456	LIB3061-050-Q1-K1-G10	LIB3061	g886470	BLASTN	1285	1e-98	82
992	456	LIB3067-043-Q1-K1-F9	LIB3067	g2738247	BLASTN	1287	1e-98	79
993	456	700087169H1	SATMON011	g1814402	BLASTN	1245	1e-94	86
994	456	LIB3069-030-Q1-K1-G9	LIB3069	g1814402	BLASTN	772	1e-93	80
995	456	LIB3069-019-Q1-K1-G11	LIB3069	g1814402	BLASTN	1202	1e-91	81
996	456	700202514H1	SATMON003	g1814402	BLASTN	1134	1e-89	84
997	456	LIB3062-011-Q1-K1-F2	LIB3062	g886470	BLASTN	664	1e-86	84
998	456	LIB143-015-Q1-E1-A1	LIB143	g974781	BLASTN	1143	1e-86	81
999	456	700570326H1	SATMON030	g974781	BLASTN	1089	1e-85	83
1000	456	700103727H1	SATMON010	g2738247	BLASTN	1132	1e-85	86
1001	456	700574823H1	SATMON030	g1814402	BLASTN	1135	1e-85	80
1002	456	700091666H1	SATMON011	g1814402	BLASTN	1136	1e-85	84
1003	456	LIB3069-033-Q1-K1-E9	LIB3069	g2738247	BLASTN	1115	1e-84	79
1004	456	700572634H1	SATMON030	g1814402	BLASTN	1123	1e-84	85
1005	456	700211829H1	SATMON016	g2738247	BLASTN	1096	1e-82	84
1006	456	700087109H1	SATMON011	g1814402	BLASTN	1102	1e-82	85
1007	456	LIB3059-003-Q1-K1-B1	LIB3059	g1814402	BLASTN	961	1e-81	82
1008	456	700047556H1	SATMON003	g2738247	BLASTN	976	1e-81	82
1009	456	700201925H1	SATMON003	g2738247	BLASTN	1001	1e-81	84
1010	456	700209343H1	SATMON016	g1814402	BLASTN	1081	1e-81	82
1011	456	700348416H1	SATMON023	g1814402	BLASTN	1081	1e-81	83
1012	456	700095559H1	SATMON008	g1814402	BLASTN	1085	1e-81	82
1013	456	700073472H1	SATMON007	g1814402	BLASTN	1089	1e-81	82
1014	456	LIB3062-057-Q1-K1-C2	LIB3062	g886470	BLASTN	1073	1e-80	78
1015	456	700093445H1	SATMON008	g1814402	BLASTN	1076	1e-80	84
1016	456	LIB143-026-Q1-E1-H3	LIB143	g2738247	BLASTN	648	1e-79	79
1017	456	700206240H1	SATMON003	g1814402	BLASTN	1057	1e-79	82
1018	456	700091907H1	SATMON011	g886470	BLASTN	1062	1e-79	80
1019	456	700258178H1	SATMON017	g1814402	BLASTN	1064	1e-79	84
1020	456	700086565H1	SATMON011	g1814402	BLASTN	1065	1e-79	82
1021	456	700471452H1	SATMON025	g2738247	BLASTN	1045	1e-78	85
1022	456	700243858H1	SATMON010	g1814402	BLASTN	1048	1e-78	85
1023	456	700352224H1	SATMON023	g2738247	BLASTN	1051	1e-78	84
1024	456	700084769H1	SATMON011	g1814402	BLASTN	1054	1e-78	83
1025	456	700082818H1	SATMON011	g886470	BLASTN	984	1e-77	84
1026	456	700331974H1	SATMON019	g886470	BLASTN	1031	1e-77	82
1027	456	700086203H1	SATMON011	g1814402	BLASTN	1032	1e-77	81
1028	456	700075826H1	SATMON007	g2738247	BLASTN	1038	1e-77	82
1029	456	700104555H1	SATMON010	g886470	BLASTN	1039	1e-77	79
1030	456	700074084H1	SATMON007	g2738247	BLASTN	1039	1e-77	87
1031	456	700076872H1	SATMON007	g1814402	BLASTN	700	1e-76	83
1032	456	700050704H1	SATMON003	g1814402	BLASTN	880	1e-76	82
1033	456	700030215H1	SATMON003	g974781	BLASTN	1019	1e-76	83
1034	456	700077416H1	SATMON007	g886470	BLASTN	1019	1e-76	79
1035	456	700093982H1	SATMON008	g1814402	BLASTN	1019	1e-76	82
1036	456	700048847H1	SATMON003	g2738247	BLASTN	1023	1e-76	85
1037	456	700090739H1	SATMON011	g886470	BLASTN	1026	1e-76	82
1038	456	700092904H1	SATMON008	g886470	BLASTN	1026	1e-76	82

1039	456	LIB3068-009-Q1-K1-E5	LIB3068	g1814402	BLASTN	1027	1e-76	73
1040	456	700336705H1	SATMON019	g1814402	BLASTN	747	1e-75	83
1041	456	700344872H1	SATMON021	g1814402	BLASTN	768	1e-75	85
1042	456	LIB3066-019-Q1-K1-E1	LIB3066	g886470	BLASTN	868	1e-75	81
1043	456	700086322H1	SATMON011	g2738247	BLASTN	1015	1e-75	81
1044	456	LIB3067-052-Q1-K1-B12	LIB3067	g974781	BLASTN	1033	1e-75	81
1045	456	700221395H1	SATMON011	g2738247	BLASTN	1000	1e-74	85
1046	456	700092483H1	SATMON008	g1814402	BLASTN	1001	1e-74	82
1047	456	700618708H1	SATMON034	g886470	BLASTN	1003	1e-74	82
1048	456	700074191H1	SATMON007	g2738247	BLASTN	1005	1e-74	86
1049	456	LIB143-055-Q1-E1-F10	LIB143	g1814402	BLASTN	986	1e-73	83
1050	456	700025955H1	SATMON003	g1814402	BLASTN	993	1e-73	84
1051	456	700574824H1	SATMON030	g886470	BLASTN	564	1e-72	80
1052	456	700201527H1	SATMON003	g974781	BLASTN	812	1e-72	81
1053	456	700092202H1	SATMON008	g886470	BLASTN	946	1e-72	79
1054	456	700223674H1	SATMON011	g2738247	BLASTN	971	1e-72	83
1055	456	700092638H1	SATMON008	g1814402	BLASTN	972	1e-72	84
1056	456	700090025H1	SATMON011	g974781	BLASTN	973	1e-72	81
1057	456	700217069H1	SATMON016	g2738247	BLASTN	976	1e-72	83
1058	456	700349142H1	SATMON023	g1814402	BLASTN	980	1e-72	82
1059	456	700088387H1	SATMON011	g2738247	BLASTN	980	1e-72	85
1060	456	700215102H1	SATMON016	g886470	BLASTN	980	1e-72	79
1061	456	700456708H1	SATMON029	g2738247	BLASTN	982	1e-72	82
1062	456	700210025H1	SATMON016	g1814402	BLASTN	982	1e-72	81
1063	456	700335345H1	SATMON019	g974781	BLASTN	982	1e-72	84
1064	456	700085257H1	SATMON011	g1814402	BLASTN	554	1e-71	82
1065	456	700026781H1	SATMON003	g974781	BLASTN	961	1e-71	82
1066	456	700224265H1	SATMON011	g2738247	BLASTN	962	1e-71	83
1067	456	700083282H1	SATMON011	g1814402	BLASTN	962	1e-71	83
1068	456	700381324H1	SATMON023	g1814402	BLASTN	965	1e-71	82
1069	456	700212017H1	SATMON016	g1814402	BLASTN	640	1e-70	83
1070	456	700088053H1	SATMON011	g886470	BLASTN	853	1e-70	84
1071	456	LIB3062-032-Q1-K1-C2	LIB3062	g1814402	BLASTN	883	1e-70	80
1072	456	700083030H1	SATMON011	g1814402	BLASTN	948	1e-70	81
1073	456	700613980H1	SATMON033	g1814402	BLASTN	950	1e-70	83
1074	456	700347019H1	SATMON021	g974781	BLASTN	952	1e-70	81
1075	456	700335621H1	SATMON019	g1814402	BLASTN	952	1e-70	80
1076	456	700094101H1	SATMON008	g1814402	BLASTN	953	1e-70	83
1077	456	700073128H1	SATMON007	g2738247	BLASTN	954	1e-70	78
1078	456	700076245H1	SATMON007	g1814402	BLASTN	955	1e-70	83
1079	456	700071903H1	SATMON007	g886470	BLASTN	956	1e-70	81
1080	456	700090035H1	SATMON011	g886470	BLASTN	935	1e-69	80
1081	456	700405323H1	SATMON029	g1814402	BLASTN	937	1e-69	83
1082	456	700050121H1	SATMON003	g886470	BLASTN	938	1e-69	83
1083	456	700224714H1	SATMON011	g1814402	BLASTN	939	1e-69	82
1084	456	700281535H2	SATMON019	g1814402	BLASTN	940	1e-69	81
1085	456	700222093H1	SATMON011	g2738247	BLASTN	941	1e-69	81
1086	456	700094659H1	SATMON008	g1814402	BLASTN	942	1e-69	84
1087	456	700237187H1	SATMON010	g1814402	BLASTN	942	1e-69	84
1088	456	700023210H1	SATMON003	g1814402	BLASTN	943	1e-69	83
1089	456	700072813H1	SATMON007	g1814402	BLASTN	944	1e-69	82
1090	456	700075720H1	SATMON007	g1814402	BLASTN	946	1e-69	83
1091	456	700072107H1	SATMON007	g1814402	BLASTN	946	1e-69	80
1092	456	700082225H1	SATMON011	g1814402	BLASTN	691	1e-68	82



1147	456	700093168H1	SATMON008	g1814402	BLASTN	885	1e-64	86
1148	456	700224478H1	SATMON011	g886470	BLASTN	886	1e-64	81
1149	456	700261958H1	SATMON017	g1814402	BLASTN	781	1e-63	79
1150	456	700074221H1	SATMON007	g1814402	BLASTN	825	1e-63	81
1151	456	700089966H1	SATMON011	g1814402	BLASTN	863	1e-63	84
1152	456	700210422H1	SATMON016	g1814402	BLASTN	863	1e-63	81
1153	456	700160272H1	SATMON012	g1814402	BLASTN	863	1e-63	85
1154	456	700444282H1	SATMON027	g2738247	BLASTN	549	1e-62	84
1155	456	700206489H1	SATMON003	g2738247	BLASTN	597	1e-62	80
1156	456	700446452H1	SATMON027	g974781	BLASTN	711	1e-62	80
1157	456	700265473H1	SATMON017	g1814402	BLASTN	715	1e-62	79
1158	456	700473708H1	SATMON025	g1814402	BLASTN	762	1e-62	82
1159	456	LIB3069-025-Q1-K1-A7	LIB3069	g886470	BLASTN	852	1e-62	80
1160	456	700166631H1	SATMON013	g2738247	BLASTN	858	1e-62	85
1161	456	700582413H1	SATMON031	g1814402	BLASTN	861	1e-62	78
1162	456	700071904H1	SATMON007	g2738247	BLASTN	544	1e-61	77
1163	456	700622655H1	SATMON034	g1814402	BLASTN	592	1e-61	82
1164	456	700082541H1	SATMON011	g1814402	BLASTN	793	1e-61	82
1165	456	700335378H1	SATMON019	g2738247	BLASTN	841	1e-61	76
1166	456	700203585H1	SATMON003	g2738247	BLASTN	846	1e-61	86
1167	456	700053043H1	SATMON007	g2738247	BLASTN	847	1e-61	81
1168	456	700163654H1	SATMON013	g1814402	BLASTN	474	1e-60	84
1169	456	700352269H1	SATMON023	g1814402	BLASTN	479	1e-60	84
1170	456	700106187H1	SATMON010	g1814402	BLASTN	631	1e-60	81
1171	456	LIB143-007-Q1-E1-A7	LIB143	g886470	BLASTN	759	1e-60	76
1172	456	700458694H1	SATMON029	g1814402	BLASTN	827	1e-60	86
1173	456	700213509H1	SATMON016	g886470	BLASTN	834	1e-60	81
1174	456	700150191H1	SATMON007	g2738247	BLASTN	834	1e-60	82
1175	456	700157594H1	SATMON012	g1814402	BLASTN	835	1e-60	80
1176	456	700094046H1	SATMON008	g974781	BLASTN	836	1e-60	79
1177	456	700220254H1	SATMON011	g2738247	BLASTN	687	1e-59	77
1178	456	700095485H1	SATMON008	g886470	BLASTN	819	1e-59	77
1179	456	LIB143-006-Q1-E1-B3	LIB143	g1814402	BLASTN	830	1e-59	65
1180	456	700071763H1	SATMON007	g2738247	BLASTN	565	1e-58	75
1181	456	700082259H1	SATMON011	g1814402	BLASTN	707	1e-58	84
1182	456	700020717H1	SATMON001	g886470	BLASTN	805	1e-58	81
1183	456	700041892H1	SATMON004	g1814402	BLASTN	805	1e-58	86
1184	456	700356573H1	SATMON024	g1814402	BLASTN	811	1e-58	81
1185	456	700171632H1	SATMON013	g2738247	BLASTN	811	1e-58	85
1186	456	700222516H1	SATMON011	g1814402	BLASTN	812	1e-58	87
1187	456	700466679H1	SATMON025	g1814402	BLASTN	812	1e-58	87
1188	456	700208929H1	SATMON016	g1814402	BLASTN	407	1e-57	78
1189	456	700355014H1	SATMON024	g1814402	BLASTN	592	1e-57	85
1190	456	701159834H1	SATMONN04	g1814402	BLASTN	683	1e-57	77
1191	456	700351137H1	SATMON023	g974781	BLASTN	725	1e-57	79
1192	456	700027308H1	SATMON003	g2738247	BLASTN	791	1e-57	80
1193	456	700142587H1	SATMON012	g974781	BLASTN	791	1e-57	83
1194	456	700170157H1	SATMON013	g1814402	BLASTN	791	1e-57	81
1195	456	700019438H1	SATMON001	g974781	BLASTN	798	1e-57	82
1196	456	700166430H1	SATMON013	g886470	BLASTN	799	1e-57	84
1197	456	700047369H1	SATMON003	g886470	BLASTN	799	1e-57	80
1198	456	700152373H1	SATMON007	g2738247	BLASTN	799	1e-57	84
1199	456	700050824H1	SATMON003	g2738247	BLASTN	605	1e-56	81
1200	456	700204588H1	SATMON003	g886470	BLASTN	720	1e-56	82

1201	456	700071667H1	SATMON007	g1814402	BLASTN	787	1e-56	85
1202	456	701185269H1	SATMONN06	g886470	BLASTN	598	1e-55	76
1203	456	700243844H1	SATMON010	g2738247	BLASTN	769	1e-55	79
1204	456	700221969H1	SATMON011	g974781	BLASTN	771	1e-55	79
1205	456	700335706H1	SATMON019	g1814402	BLASTN	773	1e-55	80
1206	456	700622285H1	SATMON034	g1814402	BLASTN	774	1e-55	82
1207	456	700089665H1	SATMON011	g2738247	BLASTN	774	1e-55	78
1208	456	700464833H1	SATMON025	g2738247	BLASTN	455	1e-54	72
1209	456	LIB3069-057-Q1-K1-A6	LIB3069	g974781	BLASTN	564	1e-54	76
1210	456	700156818H1	SATMON012	g974781	BLASTN	757	1e-54	81
1211	456	700219420H1	SATMON011	g974781	BLASTN	758	1e-54	79
1212	456	700152570H1	SATMON007	g1814402	BLASTN	761	1e-54	86
1213	456	700084484H1	SATMON011	g1814402	BLASTN	764	1e-54	84
1214	456	700152208H1	SATMON007	g1814402	BLASTN	765	1e-54	82
1215	456	700550889H1	SATMON022	g2738247	BLASTN	546	1e-53	76
1216	456	700025869H1	SATMON003	g2738247	BLASTN	675	1e-53	79
1217	456	700383075H1	SATMON024	g974781	BLASTN	724	1e-53	81
1218	456	700575104H1	SATMON030	g1814402	BLASTN	745	1e-53	70
1219	456	LIB3069-022-Q1-K1-C1	LIB3069	g974781	BLASTN	768	1e-53	81
1220	456	700215272H1	SATMON016	g1814402	BLASTN	674	1e-52	83
1221	456	700167881H1	SATMON013	g974781	BLASTN	737	1e-52	79
1222	456	700019727H1	SATMON001	g2738247	BLASTN	737	1e-52	81
1223	456	700351776H1	SATMON023	g886470	BLASTN	741	1e-52	76
1224	456	700210543H1	SATMON016	g974781	BLASTN	742	1e-52	85
1225	456	700571604H1	SATMON030	g1814402	BLASTN	360	1e-51	76
1226	456	700169777H1	SATMON013	g1814402	BLASTN	426	1e-51	83
1227	456	700096744H1	SATMON008	g2738247	BLASTN	539	1e-51	79
1228	456	700383157H1	SATMON024	g974781	BLASTN	667	1e-51	79
1229	456	700155764H1	SATMON007	g886470	BLASTN	721	1e-51	75
1230	456	700150817H1	SATMON007	g1814402	BLASTN	727	1e-51	75
1231	456	700619660H1	SATMON034	g886470	BLASTN	727	1e-51	73
1232	456	700471085H1	SATMON025	g1814402	BLASTN	406	1e-50	79
1233	456	700163226H1	SATMON013	g2738247	BLASTN	475	1e-50	78
1234	456	700444567H1	SATMON027	g1814402	BLASTN	498	1e-50	81
1235	456	700457303H1	SATMON029	g2738247	BLASTN	662	1e-50	80
1236	456	700088065H1	SATMON011	g886470	BLASTN	707	1e-50	79
1237	456	700152592H1	SATMON007	g886470	BLASTN	716	1e-50	80
1238	456	LIB3068-061-Q1-K1-B2	LIB3068	g886470	BLASTN	730	1e-50	78
1239	456	700331880H1	SATMON019	g1814402	BLASTN	697	1e-49	85
1240	456	700218025H1	SATMON016	g886470	BLASTN	700	1e-49	79
1241	456	700348643H1	SATMON023	g1814402	BLASTN	701	1e-49	80
1242	456	700236975H1	SATMON010	g974781	BLASTN	704	1e-49	81
1243	456	700442968H1	SATMON026	g886470	BLASTN	418	1e-48	79
1244	456	700479529H1	SATMON034	g2738247	BLASTN	508	1e-48	75
1245	456	700160983H1	SATMON012	g1814402	BLASTN	690	1e-48	83
1246	456	700050185H1	SATMON003	g974781	BLASTN	421	1e-47	75
1247	456	700611764H1	SATMON022	g1814402	BLASTN	503	1e-47	79
1248	456	700622669H1	SATMON034	g1814402	BLASTN	549	1e-47	79
1249	456	700153237H1	SATMON007	g1814402	BLASTN	672	1e-47	80
1250	456	700242703H1	SATMON010	g886470	BLASTN	682	1e-47	81
1251	456	700165177H1	SATMON013	g2738247	BLASTN	682	1e-47	78
1252	456	700379889H1	SATMON021	g886470	BLASTN	682	1e-47	85
1253	456	700449243H1	SATMON028	g1814402	BLASTN	357	1e-46	83
1254	456	700453282H1	SATMON028	g886470	BLASTN	608	1e-46	81

1255	456	700224308H1	SATMON011	g1814402	BLASTN	661	1e-46	83
1256	456	700150484H1	SATMON007	g886470	BLASTN	663	1e-46	78
1257	456	700240609H1	SATMON010	g2738247	BLASTN	668	1e-46	79
1258	456	700171610H1	SATMON013	g974781	BLASTN	669	1e-46	79
1259	456	LIB143-027-Q1-E1-H3	LIB143	g1814402	BLASTN	670	1e-46	86
1260	456	700334084H1	SATMON019	g1814402	BLASTN	467	1e-45	81
1261	456	LIB3069-043-Q1-K1-H6	LIB3069	g886470	BLASTN	546	1e-45	83
1262	456	700222515H1	SATMON011	g1814402	BLASTN	648	1e-45	75
1263	456	700223749H1	SATMON011	g1814402	BLASTN	648	1e-45	75
1264	456	701180187H1	SATMONN05	g974781	BLASTN	648	1e-45	78
1265	456	700050111H1	SATMON003	g2738247	BLASTN	653	1e-45	82
1266	456	700051275H1	SATMON003	g1814402	BLASTN	379	1e-44	75
1267	456	700235202H1	SATMON010	g2738247	BLASTN	637	1e-44	80
1268	456	700257385H1	SATMON017	g1814402	BLASTN	638	1e-44	80
1269	456	LIB143-028-Q1-E1-H7	LIB143	g1814402	BLASTN	638	1e-44	80
1270	456	700576066H1	SATMON030	g886470	BLASTN	516	1e-43	76
1271	456	700455065H1	SATMON029	g886470	BLASTN	624	1e-43	79
1272	456	700074158H1	SATMON007	g1814402	BLASTN	630	1e-43	83
1273	456	700171322H1	SATMON013	g1814402	BLASTN	634	1e-43	77
1274	456	700457369H1	SATMON029	g974781	BLASTN	392	1e-42	83
1275	456	700378252H1	SATMON019	g2738247	BLASTN	462	1e-42	77
1276	456	700454626H1	SATMON029	g1814402	BLASTN	495	1e-42	85
1277	456	700208182H1	SATMON016	g2738247	BLASTN	564	1e-42	73
1278	456	700618848H1	SATMON034	g886470	BLASTN	615	1e-42	80
1279	456	700222969H1	SATMON011	g1814402	BLASTN	622	1e-42	74
1280	456	700444782H1	SATMON027	g1814402	BLASTN	337	1e-41	81
1281	456	700204404H1	SATMON003	g2738247	BLASTN	607	1e-41	80
1282	456	700347028H1	SATMON021	g1814402	BLASTN	598	1e-40	81
1283	456	700172447H1	SATMON013	g1814402	BLASTN	328	1e-39	85
1284	456	700549696H1	SATMON022	g886470	BLASTN	360	1e-39	81
1285	456	700257287H1	SATMON017	g1814402	BLASTN	365	1e-38	84
1286	456	700569630H1	SATMON030	g1814402	BLASTN	566	1e-38	78
1287	456	700207258H1	SATMON017	g1814402	BLASTN	568	1e-38	85
1288	456	700052467H1	SATMON003	g1814402	BLASTN	515	1e-37	82
1289	456	700083656H1	SATMON011	g2738247	BLASTN	552	1e-37	83
1290	456	700429279H1	SATMONN01	g2738247	BLASTN	556	1e-37	79
1291	456	700349921H1	SATMON023	g2738247	BLASTN	471	1e-36	78
1292	456	700449609H1	SATMON028	g886470	BLASTN	540	1e-36	81
1293	456	700150152H1	SATMON007	g1814402	BLASTN	543	1e-36	74
1294	456	700075675H1	SATMON007	g1814402	BLASTN	546	1e-36	81
1295	456	700267015H1	SATMON017	g886470	BLASTN	550	1e-36	75
1296	456	700465118H1	SATMON025	g1814402	BLASTN	555	1e-36	78
1297	456	700456662H1	SATMON029	g2738247	BLASTN	314	1e-34	76
1298	456	700405315H1	SATMON029	g886470	BLASTN	519	1e-34	83
1299	456	700218639H1	SATMON011	g974781	BLASTN	524	1e-34	81
1300	456	700216606H1	SATMON016	g1814402	BLASTN	524	1e-34	83
1301	456	700456965H1	SATMON029	g2738247	BLASTN	525	1e-34	78
1302	456	700102147H1	SATMON010	g974781	BLASTN	526	1e-34	82
1303	456	700235734H1	SATMON010	g2738247	BLASTN	526	1e-34	83
1304	456	LIB3068-009-Q1-K1-E10	LIB3068	g886470	BLASTN	548	1e-34	70
1305	456	700029363H1	SATMON003	g974781	BLASTN	377	1e-33	80
1306	456	LIB3069-030-Q1-K1-D10	LIB3069	g886470	BLASTN	297	1e-30	78
1307	456	700334895H1	SATMON019	g974781	BLASTN	468	1e-30	82
1308	456	700150963H1	SATMON007	g974781	BLASTN	459	1e-29	88



1309	456	700152817H1	SATMON007	g974781	BLASTN	459	1e-29	88
1310	456	700208732H1	SATMON016	g974781	BLASTN	485	1e-29	76
1311	456	700051461H1	SATMON003	g886470	BLASTN	467	1e-28	73
1312	456	700616589H1	SATMON033	g2738247	BLASTN	273	1e-27	78
1313	456	700236150H1	SATMON010	g974781	BLASTN	434	1e-27	84
1314	456	700453369H1	SATMON028	g1814402	BLASTN	436	1e-27	84
1315	456	700621739H1	SATMON034	g1814402	BLASTN	436	1e-27	69
1316	456	700075162H1	SATMON007	g1814402	BLASTN	438	1e-27	85
1317	456	700405040H1	SATMON027	g974781	BLASTN	438	1e-27	72
1318	456	700356893H1	SATMON024	g974781	BLASTN	458	1e-27	82
1319	456	701185493H1	SATMONN06	g886471	BLASTX	72	1e-26	97
1320	456	700206095H1	SATMON003	g974781	BLASTN	421	1e-26	83
1321	456	700083901H1	SATMON011	g1814402	BLASTN	421	1e-26	80
1322	456	LIB3062-026-Q1-K1-D7	LIB3062	g1814402	BLASTN	423	1e-24	81
1323	456	700377277H1	SATMON019	g1814402	BLASTN	357	1e-20	75
1324	456	700201214H1	SATMON003	g2738248	BLASTX	152	1e-18	77
1325	456	700025959H1	SATMON003	g886471	BLASTX	188	1e-18	94
1326	456	700259484H1	SATMON017	g1814402	BLASTN	189	1e-17	82
1327	456	700613969H1	SATMON033	g886470	BLASTN	315	1e-17	90
1328	456	700215602H1	SATMON016	g2738248	BLASTX	158	1e-16	74
1329	456	700438152H1	SATMON026	g2738247	BLASTN	258	1e-16	75
1330	456	700458654H1	SATMON029	g886470	BLASTN	330	1e-16	70
1331	456	700449563H1	SATMON028	g886471	BLASTX	163	1e-15	89
1332	456	700236490H1	SATMON010	g1814402	BLASTN	276	1e-14	89
1333	456	700456927H1	SATMON029	g1814403	BLASTX	119	1e-13	86
1334	456	701180088H1	SATMONN05	g886471	BLASTX	152	1e-13	60
1335	456	700455716H1	SATMON029	g2738248	BLASTX	111	1e-12	67
1336	456	700573520H1	SATMON030	g886471	BLASTX	82	1e-11	84
1337	456	700569938H1	SATMON030	g2738248	BLASTX	132	1e-11	91
1338	456	700551958H1	SATMON022	g2738247	BLASTN	264	1e-11	82
1339	456	700337475H1	SATMON020	g2738247	BLASTN	268	1e-11	83
1340	456	700170827H1	SATMON013	g886471	BLASTX	124	1e-10	77
1341	456	700453382H1	SATMON028	g974781	BLASTN	152	1e-10	85
1342	456	700089211H1	SATMON011	g2738247	BLASTN	245	1e-9	79
1343	456	700103155H1	SATMON010	g2738248	BLASTX	82	1e-8	85
1344	5523	LIB3062-017-Q1-K1-F4	LIB3062	g2738247	BLASTN	932	1e-84	76
1345	5523	700210708H1	SATMON016	g2738247	BLASTN	952	1e-70	79
1346	5523	700219307H1	SATMON011	g1814402	BLASTN	892	1e-65	78
1347	5523	700221188H1	SATMON011	g2738247	BLASTN	867	1e-63	81
1348	5523	700203884H1	SATMON003	g2738247	BLASTN	874	1e-63	77
1349	5523	700218549H1	SATMON011	g2738247	BLASTN	811	1e-58	77
1350	5523	700572845H2	SATMON030	g2738247	BLASTN	785	1e-56	77
1351	5523	700152362H1	SATMON007	g886470	BLASTN	742	1e-52	79
1352	5523	700152138H1	SATMON007	g2738247	BLASTN	714	1e-50	80
1353	5523	700218842H1	SATMON011	g2738247	BLASTN	700	1e-49	75
2666	-700697958	700697958H1	SOYMON015	g2738247	BLASTN	258	1e-10	83
2667	-700731277	700731277H1	SOYMON009	g886470	BLASTN	930	1e-68	83
2668	-700749261	700749261H1	SOYMON013	g1814402	BLASTN	508	1e-38	78
2669	-700787742	700787742H2	SOYMON011	g1814402	BLASTN	648	1e-50	81
2670	-700831146	700831146H1	SOYMON019	g1814402	BLASTN	476	1e-34	80
2671	-700832029	700832029H1	SOYMON019	g1814402	BLASTN	328	1e-27	78
2672	-700854481	700854481H1	SOYMON023	g1814402	BLASTN	193	1e-11	85
2673	-700873716	700873716H1	SOYMON018	g1749542	BLASTX	99	1e-15	53
2674	-700893904	700893904H1	SOYMON024	g886470	BLASTN	413	1e-25	82

2675	-700909658	700909658H1	SOYMON022	g1814402	BLASTN	251	1e-10	91
2676	-700943841	700943841H1	SOYMON024	g886470	BLASTN	549	1e-49	82
2677	-700963223	700963223H1	SOYMON022	g886470	BLASTN	649	1e-45	72
2678	-700974103	700974103H1	SOYMON005	g886470	BLASTN	763	1e-54	80
2679	-700994293	700994293H1	SOYMON011	g974782	BLASTX	97	1e-12	72
2680	-701004816	701004816H1	SOYMON019	g1814402	BLASTN	582	1e-39	84
2681	-701007125	701007125H1	SOYMON019	g1814403	BLASTX	152	1e-13	87
2682	-701008540	701008540H1	SOYMON019	g886470	BLASTN	770	1e-55	74
2683	-701037766	701037766H1	SOYMON029	g1814403	BLASTX	72	1e-10	69
2684	-701062191	701062191H1	SOYMON033	g974781	BLASTN	225	1e-9	79
2685	-701105474	701105474H1	SOYMON036	g974782	BLASTX	164	1e-15	91
2686	-GM14442	LIB3049-056-Q1-E1-B1	LIB3049	g974781	BLASTN	231	1e-10	79
2687	-GM19631	LIB3056-007-Q1-N1-G9	LIB3056	g974781	BLASTN	388	1e-37	78
2688	-GM37189	LIB3051-072-Q1-K1-B5	LIB3051	g1814402	BLASTN	316	1e-15	78
2689	-GM44802	LIB3053-004-Q1-N1-B2	LIB3053	g974782	BLASTX	85	1e-26	90
2690	1382	700683236H1	SOYMON008	g886470	BLASTN	805	1e-58	78
2691	1382	700566434H1	SOYMON002	g886471	BLASTX	162	1e-23	76
2692	15690	701064361H1	SOYMON034	g974781	BLASTN	951	1e-70	84
2693	15690	700847301H1	SOYMON021	g886470	BLASTN	707	1e-66	85
2694	17335	LIB3051-088-Q1-K1-D9	LIB3051	g886470	BLASTN	1285	1e-101	79
2695	17335	701003832H1	SOYMON019	g974781	BLASTN	811	1e-58	77
2696	17335	700864888H1	SOYMON016	g1814402	BLASTN	767	1e-55	79
2697	17335	700672491H1	SOYMON006	g974781	BLASTN	428	1e-46	75
2698	17335	701003457H1	SOYMON019	g1814402	BLASTN	411	1e-29	84
2699	17335	700833672H1	SOYMON019	g974781	BLASTN	445	1e-28	79
2700	17900	700850578H1	SOYMON023	g2738247	BLASTN	905	1e-66	82
2701	17900	701053154H1	SOYMON032	g1814402	BLASTN	725	1e-65	83
2702	17900	700842351H1	SOYMON020	g1814402	BLASTN	878	1e-64	83
2703	17900	700837656H1	SOYMON020	g1814402	BLASTN	841	1e-61	83
2704	17900	700890851H1	SOYMON024	g974781	BLASTN	747	1e-53	80
2705	20688	700908840H1	SOYMON022	g2738247	BLASTN	889	1e-65	82
2706	20688	700908848H1	SOYMON022	g2738247	BLASTN	877	1e-64	81
2707	33542	LIB3051-009-Q1-E1-E5	LIB3051	g974781	BLASTN	1218	1e-92	79
2708	33542	700748773H1	SOYMON013	g974781	BLASTN	669	1e-46	78
2709	33542	700836363H1	SOYMON020	g2738247	BLASTN	596	1e-40	80
2710	4243	701123616H1	SOYMON037	g1814402	BLASTN	995	1e-74	87
2711	4243	700555001H1	SOYMON001	g1814402	BLASTN	975	1e-72	90
2712	4243	701002967H1	SOYMON019	g1814402	BLASTN	950	1e-70	86
2713	4243	700653509H1	SOYMON003	g1814402	BLASTN	539	1e-69	85
2714	4243	701206028H1	SOYMON035	g1814402	BLASTN	938	1e-69	86
2715	4243	700962115H1	SOYMON022	g1814402	BLASTN	923	1e-68	86
2716	4243	700866243H1	SOYMON016	g1814402	BLASTN	909	1e-66	82
2717	4243	700752507H1	SOYMON014	g1814402	BLASTN	887	1e-65	85
2718	4243	701003887H1	SOYMON019	g1814402	BLASTN	863	1e-63	86
2719	4243	700556913H1	SOYMON001	g1814402	BLASTN	865	1e-63	86
2720	4243	701013549H1	SOYMON019	g1814402	BLASTN	867	1e-63	90
2721	4243	701209706H1	SOYMON035	g1814402	BLASTN	871	1e-63	90
2722	4243	701010487H1	SOYMON019	g1814402	BLASTN	529	1e-62	79
2723	4243	700548246H1	SOYMON002	g1814402	BLASTN	553	1e-62	82
2724	4243	701138219H1	SOYMON038	g1814402	BLASTN	594	1e-62	86
2725	4243	700965160H1	SOYMON022	g1814402	BLASTN	852	1e-62	91
2726	4243	701015168H1	SOYMON019	g1814402	BLASTN	855	1e-62	89
2727	4243	701136095H1	SOYMON038	g1814402	BLASTN	839	1e-61	88
2728	4243	700761789H1	SOYMON015	g1814402	BLASTN	845	1e-61	86

2729	4243	701105695H1	SOYMON036	g1814402	BLASTN	835	1e-60	87
2730	4243	700991714H1	SOYMON011	g1814402	BLASTN	502	1e-59	83
2731	4243	700564223H1	SOYMON002	g1814402	BLASTN	557	1e-59	90
2732	4243	700987384H1	SOYMON009	g1814402	BLASTN	803	1e-58	86
2733	4243	700833934H1	SOYMON019	g1814402	BLASTN	806	1e-58	89
2734	4243	700835181H1	SOYMON019	g1814402	BLASTN	806	1e-58	82
2735	4243	700737529H1	SOYMON010	g1814402	BLASTN	810	1e-58	92
2736	4243	701012851H1	SOYMON019	g1814402	BLASTN	811	1e-58	91
2737	4243	700556592H1	SOYMON001	g1814402	BLASTN	814	1e-58	88
2738	4243	700907579H1	SOYMON022	g1814402	BLASTN	781	1e-56	89
2739	4243	700961749H1	SOYMON022	g1814402	BLASTN	785	1e-56	91
2740	4243	700835239H1	SOYMON019	g1814402	BLASTN	787	1e-56	86
2741	4243	700646425H1	SOYMON013	g1814402	BLASTN	772	1e-55	89
2742	4243	701123924H1	SOYMON037	g1814402	BLASTN	775	1e-55	91
2743	4243	700957759H1	SOYMON022	g1814402	BLASTN	776	1e-55	90
2744	4243	700964425H1	SOYMON022	g1814402	BLASTN	777	1e-55	86
2745	4243	700962173H1	SOYMON022	g1814402	BLASTN	778	1e-55	91
2746	4243	701066293H1	SOYMON034	g1814402	BLASTN	759	1e-54	81
2747	4243	700986741H1	SOYMON009	g1814402	BLASTN	589	1e-53	84
2748	4243	701212514H1	SOYMON035	g1814402	BLASTN	591	1e-53	87
2749	4243	701009195H1	SOYMON019	g1814402	BLASTN	747	1e-53	91
2750	4243	701060510H1	SOYMON033	g1814402	BLASTN	748	1e-53	84
2751	4243	700848730H1	SOYMON021	g1814402	BLASTN	749	1e-53	90
2752	4243	700754870H1	SOYMON014	g1814402	BLASTN	751	1e-53	84
2753	4243	701212482H1	SOYMON035	g1814402	BLASTN	751	1e-53	84
2754	4243	700753283H1	SOYMON014	g1814402	BLASTN	752	1e-53	86
2755	4243	700738517H1	SOYMON012	g1814402	BLASTN	636	1e-52	89
2756	4243	700833978H1	SOYMON019	g1814402	BLASTN	740	1e-52	91
2757	4243	700756424H1	SOYMON014	g1814402	BLASTN	729	1e-51	82
2758	4243	701011759H1	SOYMON019	g1814402	BLASTN	729	1e-51	91
2759	4243	701010103H2	SOYMON019	g1814402	BLASTN	707	1e-50	82
2760	4243	700741392H1	SOYMON012	g1814402	BLASTN	707	1e-50	84
2761	4243	701123062H1	SOYMON037	g1814402	BLASTN	308	1e-49	88
2762	4243	701048949H1	SOYMON032	g1814402	BLASTN	502	1e-49	85
2763	4243	700834566H1	SOYMON019	g1814402	BLASTN	618	1e-49	88
2764	4243	700963965H1	SOYMON022	g1814402	BLASTN	685	1e-48	78
2765	4243	700986376H1	SOYMON009	g1814402	BLASTN	694	1e-48	84
2766	4243	701012708H1	SOYMON019	g1814402	BLASTN	521	1e-47	91
2767	4243	700746927H1	SOYMON013	g1814402	BLASTN	547	1e-46	78
2768	4243	700997448H1	SOYMON018	g1814402	BLASTN	470	1e-45	89
2769	4243	700830667H1	SOYMON019	g1814402	BLASTN	647	1e-45	86
2770	4243	700891479H1	SOYMON024	g1814402	BLASTN	654	1e-45	88
2771	4243	700562246H1	SOYMON002	g1814402	BLASTN	656	1e-45	86
2772	4243	701097325H1	SOYMON028	g1814402	BLASTN	420	1e-44	92
2773	4243	700835830H1	SOYMON019	g1814402	BLASTN	503	1e-44	86
2774	4243	700962969H1	SOYMON022	g1814402	BLASTN	515	1e-44	89
2775	4243	701102860H1	SOYMON028	g1814402	BLASTN	309	1e-42	92
2776	4243	700763506H1	SOYMON015	g1814402	BLASTN	456	1e-42	84
2777	4243	700994059H1	SOYMON011	g1814402	BLASTN	487	1e-42	87
2778	4243	700751551H1	SOYMON014	g1814402	BLASTN	568	1e-41	85
2779	4243	701108872H1	SOYMON036	g1814402	BLASTN	603	1e-41	85
2780	4243	700994053H1	SOYMON011	g1814402	BLASTN	587	1e-40	85
2781	4243	700729064H1	SOYMON009	g1814402	BLASTN	413	1e-38	90
2782	4243	700874176H1	SOYMON018	g1814402	BLASTN	468	1e-38	92

2783	4243	700742963H1	SOYMON012	g1814402	BLASTN	565	1e-38	86
2784	4243	701212923H1	SOYMON035	g1814402	BLASTN	572	1e-38	84
2785	4243	700994851H1	SOYMON011	g1814402	BLASTN	548	1e-36	76
2786	4243	701000193H1	SOYMON018	g1814402	BLASTN	296	1e-33	87
2787	4243	700756219H1	SOYMON014	g1814402	BLASTN	394	1e-32	80
2788	4243	701014751H1	SOYMON019	g1814402	BLASTN	461	1e-29	90
2789	4243	700561163H1	SOYMON002	g1814402	BLASTN	231	1e-25	89
2790	4243	700650284H1	SOYMON003	g974782	BLASTX	157	1e-14	96
2791	4243	700869218H1	SOYMON016	g1814402	BLASTN	248	1e-11	93
2792	550	LIB3028-007-Q1-B1-B6	LIB3028	g886470	BLASTN	1440	1e-111	81
2793	550	LIB3040-017-Q1-E1-E8	LIB3040	g886470	BLASTN	1260	1e-96	80
2794	550	700650656H1	SOYMON003	g1814402	BLASTN	1072	1e-93	83
2795	550	LIB3051-091-Q1-K1-C2	LIB3051	g886470	BLASTN	1124	1e-89	80
2796	550	LIB3051-072-Q1-K1-B3	LIB3051	g974781	BLASTN	1114	1e-83	82
2797	550	LIB3051-006-Q1-E1-G9	LIB3051	g974781	BLASTN	703	1e-82	81
2798	550	700563811H1	SOYMON002	g1814402	BLASTN	1071	1e-80	84
2799	550	700754104H1	SOYMON014	g974781	BLASTN	1052	1e-78	86
2800	550	701002973H1	SOYMON019	g1814402	BLASTN	1039	1e-77	85
2801	550	700986733H1	SOYMON009	g974781	BLASTN	1039	1e-77	83
2802	550	700557595H1	SOYMON001	g886470	BLASTN	1019	1e-76	83
2803	550	700563477H1	SOYMON002	g2738247	BLASTN	1024	1e-76	85
2804	550	700976112H1	SOYMON009	g886470	BLASTN	1024	1e-76	84
2805	550	701004484H1	SOYMON019	g1814402	BLASTN	1007	1e-75	84
2806	550	700889161H1	SOYMON024	g974781	BLASTN	996	1e-74	85
2807	550	700833110H1	SOYMON019	g974781	BLASTN	998	1e-74	87
2808	550	701124559H1	SOYMON037	g974781	BLASTN	1002	1e-74	86
2809	550	700729207H1	SOYMON009	g974781	BLASTN	1004	1e-74	85
2810	550	700730012H1	SOYMON009	g886470	BLASTN	987	1e-73	85
2811	550	700987128H1	SOYMON009	g2738247	BLASTN	989	1e-73	85
2812	550	700564788H1	SOYMON002	g974781	BLASTN	990	1e-73	84
2813	550	701099104H1	SOYMON028	g1814402	BLASTN	994	1e-73	86
2814	550	LIB3065-008-Q1-N1-B4	LIB3065	g2738247	BLASTN	839	1e-72	83
2815	550	701104283H1	SOYMON036	g974781	BLASTN	976	1e-72	83
2816	550	700726387H1	SOYMON009	g974781	BLASTN	980	1e-72	84
2817	550	700900884H1	SOYMON027	g1814402	BLASTN	982	1e-72	85
2818	550	LIB3051-029-Q1-K1-D6	LIB3051	g886470	BLASTN	824	1e-71	81
2819	550	701213995H1	SOYMON035	g1814402	BLASTN	962	1e-71	84
2820	550	700683596H1	SOYMON008	g1814402	BLASTN	968	1e-71	85
2821	550	700646529H1	SOYMON014	g1814402	BLASTN	867	1e-70	84
2822	550	700756704H1	SOYMON014	g1814402	BLASTN	893	1e-70	87
2823	550	700991647H1	SOYMON011	g1814402	BLASTN	947	1e-70	84
2824	550	700962195H1	SOYMON022	g974781	BLASTN	947	1e-70	87
2825	550	700994369H1	SOYMON011	g1814402	BLASTN	949	1e-70	84
2826	550	700672477H1	SOYMON006	g1814402	BLASTN	951	1e-70	85
2827	550	700946215H1	SOYMON024	g2738247	BLASTN	957	1e-70	83
2828	550	701152765H1	SOYMON031	g974781	BLASTN	958	1e-70	88
2829	550	701010552H1	SOYMON019	g974781	BLASTN	501	1e-69	83
2830	550	LIB3056-004-Q1-N1-B12	LIB3056	g2738247	BLASTN	698	1e-69	77
2831	550	701100656H1	SOYMON028	g974781	BLASTN	890	1e-69	85
2832	550	700985362H1	SOYMON009	g1814402	BLASTN	937	1e-69	82
2833	550	700746178H1	SOYMON013	g974781	BLASTN	938	1e-69	85
2834	550	700674440H1	SOYMON007	g974781	BLASTN	938	1e-69	83
2835	550	700556934H1	SOYMON001	g1814402	BLASTN	938	1e-69	82
2836	550	700652624H1	SOYMON003	g1814402	BLASTN	939	1e-69	82

2837	550	700952331H1	SOYMON022	g886470	BLASTN	946	1e-69	83
2838	550	700725576H1	SOYMON009	g886470	BLASTN	946	1e-69	85
2839	550	701003602H1	SOYMON019	g2738247	BLASTN	924	1e-68	83
2840	550	700745053H1	SOYMON013	g974781	BLASTN	924	1e-68	85
2841	550	700895781H1	SOYMON027	g1814402	BLASTN	926	1e-68	84
2842	550	700664593H1	SOYMON005	g974781	BLASTN	926	1e-68	87
2843	550	700864264H1	SOYMON016	g1814402	BLASTN	930	1e-68	84
2844	550	700674466H1	SOYMON007	g974781	BLASTN	933	1e-68	83
2845	550	700564170H1	SOYMON002	g974781	BLASTN	861	1e-67	83
2846	550	700654531H1	SOYMON004	g974781	BLASTN	911	1e-67	81
2847	550	700996341H1	SOYMON018	g886470	BLASTN	911	1e-67	83
2848	550	701136257H1	SOYMON038	g886470	BLASTN	912	1e-67	82
2849	550	700751114H1	SOYMON014	g1814402	BLASTN	914	1e-67	84
2850	550	700657606H1	SOYMON004	g1814402	BLASTN	916	1e-67	87
2851	550	700983827H1	SOYMON009	g974781	BLASTN	918	1e-67	85
2852	550	700981249H1	SOYMON009	g1814402	BLASTN	921	1e-67	80
2853	550	700836103H1	SOYMON019	g886470	BLASTN	922	1e-67	82
2854	550	701011869H1	SOYMON019	g886470	BLASTN	486	1e-66	86
2855	550	701097045H1	SOYMON028	g886470	BLASTN	601	1e-66	85
2856	550	701012695H1	SOYMON019	g974781	BLASTN	777	1e-66	85
2857	550	700945396H1	SOYMON024	g974781	BLASTN	902	1e-66	85
2858	550	700755390H1	SOYMON014	g974781	BLASTN	903	1e-66	86
2859	550	700967721H1	SOYMON033	g886470	BLASTN	910	1e-66	82
2860	550	700750610H1	SOYMON014	g974781	BLASTN	910	1e-66	84
2861	550	700908231H1	SOYMON022	g974781	BLASTN	527	1e-65	83
2862	550	701003835H1	SOYMON019	g886470	BLASTN	723	1e-65	82
2863	550	700790802H1	SOYMON011	g2738247	BLASTN	891	1e-65	82
2864	550	701053438H1	SOYMON032	g974781	BLASTN	893	1e-65	82
2865	550	701009430H1	SOYMON019	g974781	BLASTN	894	1e-65	85
2866	550	700891415H1	SOYMON024	g1814402	BLASTN	895	1e-65	84
2867	550	701100681H1	SOYMON028	g2738247	BLASTN	482	1e-64	84
2868	550	700893420H1	SOYMON024	g974781	BLASTN	570	1e-64	89
2869	550	700752741H1	SOYMON014	g886470	BLASTN	880	1e-64	82
2870	550	700955938H1	SOYMON022	g1814402	BLASTN	882	1e-64	81
2871	550	701015042H1	SOYMON019	g2738247	BLASTN	886	1e-64	82
2872	550	LIB3051-072-Q1-K1-B1	LIB3051	g974781	BLASTN	645	1e-63	76
2873	550	700741918H1	SOYMON012	g886470	BLASTN	722	1e-63	84
2874	550	701103319H1	SOYMON028	g974781	BLASTN	792	1e-63	82
2875	550	700833218H1	SOYMON019	g2738247	BLASTN	863	1e-63	81
2876	550	701008071H1	SOYMON019	g886470	BLASTN	867	1e-63	83
2877	550	700832073H1	SOYMON019	g974781	BLASTN	871	1e-63	83
2878	550	700889695H1	SOYMON024	g974781	BLASTN	872	1e-63	83
2879	550	701007489H2	SOYMON019	g974781	BLASTN	873	1e-63	84
2880	550	700895858H1	SOYMON027	g974781	BLASTN	874	1e-63	84
2881	550	700753955H1	SOYMON014	g974781	BLASTN	470	1e-62	87
2882	550	700564433H1	SOYMON002	g886470	BLASTN	757	1e-62	84
2883	550	700963115H1	SOYMON022	g974781	BLASTN	851	1e-62	83
2884	550	700894728H1	SOYMON024	g886470	BLASTN	860	1e-62	84
2885	550	701056915H1	SOYMON033	g886470	BLASTN	862	1e-62	84
2886	550	700741134H1	SOYMON012	g886470	BLASTN	862	1e-62	83
2887	550	700847591H1	SOYMON021	g974781	BLASTN	842	1e-61	84
2888	550	700941253H1	SOYMON024	g2738247	BLASTN	844	1e-61	80
2889	550	701004315H1	SOYMON019	g2738247	BLASTN	846	1e-61	83
2890	550	700895720H1	SOYMON027	g974781	BLASTN	848	1e-61	83

2891	550	701013541H1	SOYMON019	g974781	BLASTN	849	1e-61	84
2892	550	700892552H1	SOYMON024	g886470	BLASTN	719	1e-60	83
2893	550	701141313H1	SOYMON038	g974781	BLASTN	827	1e-60	83
2894	550	701012547H1	SOYMON019	g1814402	BLASTN	831	1e-60	83
2895	550	701008558H1	SOYMON019	g1814402	BLASTN	831	1e-60	83
2896	550	700902022H1	SOYMON027	g2738247	BLASTN	831	1e-60	82
2897	550	700959515H1	SOYMON022	g886470	BLASTN	832	1e-60	81
2898	550	701042630H1	SOYMON029	g1814402	BLASTN	819	1e-59	81
2899	550	700941292H1	SOYMON024	g2738247	BLASTN	820	1e-59	81
2900	550	700788526H1	SOYMON011	g974781	BLASTN	491	1e-58	82
2901	550	700894839H1	SOYMON024	g1814402	BLASTN	498	1e-58	82
2902	550	700865873H1	SOYMON016	g974781	BLASTN	808	1e-58	85
2903	550	701015435H1	SOYMON019	g886470	BLASTN	809	1e-58	80
2904	550	700755960H1	SOYMON014	g2738247	BLASTN	809	1e-58	78
2905	550	700876051H1	SOYMON018	g886470	BLASTN	434	1e-57	81
2906	550	701041327H1	SOYMON029	g886470	BLASTN	499	1e-57	83
2907	550	701098902H1	SOYMON028	g2738247	BLASTN	767	1e-57	77
2908	550	700853392H1	SOYMON023	g886470	BLASTN	793	1e-57	82
2909	550	700872645H1	SOYMON018	g974781	BLASTN	798	1e-57	83
2910	550	700989675H1	SOYMON011	g2738247	BLASTN	800	1e-57	79
2911	550	700753487H1	SOYMON014	g1814402	BLASTN	589	1e-56	83
2912	550	700736276H1	SOYMON010	g1814402	BLASTN	782	1e-56	79
2913	550	700891361H1	SOYMON024	g1814402	BLASTN	783	1e-56	81
2914	550	700829712H1	SOYMON019	g974781	BLASTN	790	1e-56	80
2915	550	LIB3050-019-Q1-K1-A1	LIB3050	g974781	BLASTN	663	1e-55	81
2916	550	701001013H1	SOYMON018	g1814402	BLASTN	768	1e-55	84
2917	550	LIB3028-031-Q1-B1-G12	LIB3028	g886470	BLASTN	775	1e-55	82
2918	550	701212782H1	SOYMON035	g886470	BLASTN	621	1e-54	82
2919	550	701008695H1	SOYMON019	g974781	BLASTN	718	1e-54	80
2920	550	700990972H1	SOYMON011	g1814402	BLASTN	766	1e-54	81
2921	550	700789576H2	SOYMON011	g886470	BLASTN	766	1e-54	80
2922	550	700994266H1	SOYMON011	g1814402	BLASTN	408	1e-53	85
2923	550	700731985H1	SOYMON010	g974781	BLASTN	590	1e-53	81
2924	550	700907927H1	SOYMON022	g886470	BLASTN	612	1e-53	80
2925	550	701012079H1	SOYMON019	g974781	BLASTN	669	1e-53	86
2926	550	700753939H1	SOYMON014	g886470	BLASTN	690	1e-53	78
2927	550	701000754H1	SOYMON018	g974781	BLASTN	745	1e-53	76
2928	550	701040287H1	SOYMON029	g886470	BLASTN	747	1e-53	82
2929	550	700891329H1	SOYMON024	g974781	BLASTN	749	1e-53	79
2930	550	700897258H1	SOYMON027	g886470	BLASTN	752	1e-53	86
2931	550	700944949H1	SOYMON024	g974781	BLASTN	426	1e-52	82
2932	550	701108671H1	SOYMON036	g1814402	BLASTN	731	1e-52	77
2933	550	700905233H1	SOYMON022	g886470	BLASTN	732	1e-52	77
2934	550	700958589H1	SOYMON022	g886470	BLASTN	738	1e-52	80
2935	550	700666436H1	SOYMON005	g2738247	BLASTN	739	1e-52	82
2936	550	700829902H1	SOYMON019	g886470	BLASTN	740	1e-52	82
2937	550	700740110H1	SOYMON012	g974781	BLASTN	742	1e-52	83
2938	550	700989055H1	SOYMON011	g886470	BLASTN	538	1e-50	79
2939	550	701213370H1	SOYMON035	g974781	BLASTN	599	1e-50	83
2940	550	701098072H1	SOYMON028	g974781	BLASTN	709	1e-50	74
2941	550	700896128H1	SOYMON027	g886470	BLASTN	714	1e-50	83
2942	550	701060755H1	SOYMON033	g974781	BLASTN	716	1e-50	74
2943	550	700953594H1	SOYMON022	g1814402	BLASTN	695	1e-49	78
2944	550	701046911H1	SOYMON032	g2738247	BLASTN	687	1e-48	81

2945	550	701065707H1	SOYMON034	g1814402	BLASTN	443	1e-47	84
2946	550	700962114H1	SOYMON022	g886470	BLASTN	678	1e-47	84
2947	550	700831826H1	SOYMON019	g2738247	BLASTN	682	1e-47	77
2948	550	701054296H1	SOYMON032	g886470	BLASTN	454	1e-46	83
2949	550	700888738H1	SOYMON024	g974781	BLASTN	552	1e-46	77
2950	550	700892022H1	SOYMON024	g886470	BLASTN	605	1e-46	81
2951	550	700890275H1	SOYMON024	g2738247	BLASTN	666	1e-46	77
2952	550	LIB3051-006-Q1-K1-G9	LIB3051	g974781	BLASTN	670	1e-45	80
2953	550	700889113H1	SOYMON024	g886470	BLASTN	582	1e-43	81
2954	550	700952720H1	SOYMON022	g886470	BLASTN	611	1e-42	76
2955	550	701014761H1	SOYMON019	g886470	BLASTN	318	1e-41	84
2956	550	700753882H1	SOYMON014	g886470	BLASTN	381	1e-41	79
2957	550	700743792H1	SOYMON012	g1814402	BLASTN	610	1e-41	86
2958	550	700990963H1	SOYMON011	g886470	BLASTN	578	1e-39	77
2959	550	700941880H1	SOYMON024	g2738247	BLASTN	583	1e-39	81
2960	550	700898962H1	SOYMON027	g2738247	BLASTN	571	1e-38	80
2961	550	700990865H1	SOYMON011	g2738247	BLASTN	467	1e-37	77
2962	550	700565779H1	SOYMON002	g886470	BLASTN	557	1e-37	70
2963	550	700993903H1	SOYMON011	g974781	BLASTN	562	1e-37	84
2964	550	700941589H1	SOYMON024	g2738247	BLASTN	550	1e-36	81
2965	550	701052554H1	SOYMON032	g886470	BLASTN	534	1e-35	67
2966	550	700991055H1	SOYMON011	g886470	BLASTN	356	1e-34	79
2967	550	701010438H1	SOYMON019	g2738247	BLASTN	508	1e-33	79
2968	550	700756634H1	SOYMON014	g2738247	BLASTN	494	1e-32	76
2969	550	701042980H1	SOYMON029	g886470	BLASTN	461	1e-29	83
2970	550	700682940H1	SOYMON008	g2738247	BLASTN	466	1e-29	83
2971	550	701049575H1	SOYMON032	g886470	BLASTN	433	1e-27	82
2972	550	700982552H1	SOYMON009	g886470	BLASTN	356	1e-25	78
2973	550	700675637H1	SOYMON007	g886470	BLASTN	375	1e-25	79
2974	550	701142153H1	SOYMON038	g886470	BLASTN	377	1e-22	76
2975	550	700682724H1	SOYMON008	g974781	BLASTN	361	1e-19	88
2976	550	701051764H1	SOYMON032	g1814402	BLASTN	211	1e-17	80
2977	550	700867241H1	SOYMON016	g2738248	BLASTX	152	1e-13	88
2978	550	701054954H1	SOYMON032	g2738248	BLASTX	138	1e-12	86
2979	550	700790450H2	SOYMON011	g974781	BLASTN	238	1e-10	80
2980	550	700653979H1	SOYMON003	g1814403	BLASTX	118	1e-9	92
2981	550	700894218H1	SOYMON024	g2738248	BLASTX	122	1e-9	78
2982	550	700863078H1	SOYMON022	g2738247	BLASTN	236	1e-8	78
2983	5758	701209304H1	SOYMON035	g886470	BLASTN	766	1e-54	84
2984	5758	701106455H1	SOYMON036	g886470	BLASTN	723	1e-51	82
2985	5758	700833538H1	SOYMON019	g1814402	BLASTN	609	1e-43	83
2986	5758	701051425H1	SOYMON032	g886470	BLASTN	629	1e-43	83
2987	5758	700654506H1	SOYMON004	g886470	BLASTN	438	1e-26	70
2988	5758	701047795H1	SOYMON032	g974782	BLASTX	161	1e-15	100
2989	5758	701202409H1	SOYMON035	g1814402	BLASTN	310	1e-15	79
2990	8266	700558628H1	SOYMON001	g886470	BLASTN	780	1e-56	74
2991	8266	701207720H1	SOYMON035	g1814402	BLASTN	766	1e-54	74
2992	8266	700557429H1	SOYMON001	g1814402	BLASTN	728	1e-51	75

#### ADENOSYLHOMOCYSTEINASE ( EC 3.3.1.1 )

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1354	-700154280	700154280H1	SATMON007	g170772	BLASTN	229	1e-23	77

1355	-L30594291	LIB3059-032-Q1-K1-A8	LIB3059	g170772	BLASTN	516	1e-74	73
1356	-L30664307	LIB3066-047-Q1-K1-E7	LIB3066	g170772	BLASTN	628	1e-49	70
1357	503	LIB3079-013-Q1-K1-D3	LIB3079	g170772	BLASTN	1505	1e-139	89
1358	503	LIB3062-017-Q1-K1-A10	LIB3062	g170772	BLASTN	1571	1e-129	89
1359	503	LIB3067-001-Q1-K1-D11	LIB3067	g170772	BLASTN	1654	1e-129	89
1360	503	LIB148-060-Q1-E1-B9	LIB148	g170772	BLASTN	1620	1e-126	90
1361	503	LIB3069-043-Q1-K1-G4	LIB3069	g170772	BLASTN	1360	1e-124	84
1362	503	LIB143-006-Q1-E1-F3	LIB143	g170772	BLASTN	831	1e-120	88
1363	503	LIB189-009-Q1-E1-E9	LIB189	g170772	BLASTN	1551	1e-120	90
1364	503	LIB3060-003-Q1-K1-F9	LIB3060	g170772	BLASTN	1538	1e-119	91
1365	503	700089023H1	SATMON011	g170772	BLASTN	1495	1e-115	91
1366	503	LIB143-061-Q1-E1-C9	LIB143	g170772	BLASTN	1474	1e-114	90
1367	503	LIB3069-034-Q1-K1-E5	LIB3069	g170772	BLASTN	1483	1e-114	88
1368	503	LIB143-061-Q1-E1-E5	LIB143	g170772	BLASTN	1440	1e-113	89
1369	503	LIB3067-040-Q1-K1-H5	LIB3067	g170772	BLASTN	1467	1e-113	88
1370	503	LIB3067-048-Q1-K1-A11	LIB3067	g170772	BLASTN	1207	1e-109	86
1371	503	LIB3068-050-Q1-K1-G6	LIB3068	g170772	BLASTN	1279	1e-109	86
1372	503	LIB3069-036-Q1-K1-F6	LIB3069	g170772	BLASTN	1310	1e-109	92
1373	503	LIB3066-047-Q1-K1-H5	LIB3066	g170772	BLASTN	1353	1e-108	86
1374	503	LIB3059-011-Q1-K1-A5	LIB3059	g170772	BLASTN	1401	1e-107	90
1375	503	LIB3067-056-Q1-K1-E11	LIB3067	g170772	BLASTN	957	1e-106	86
1376	503	LIB189-010-Q1-E1-E12	LIB189	g170772	BLASTN	1144	1e-106	88
1377	503	700084426H1	SATMON011	g170772	BLASTN	1368	1e-105	91
1378	503	700086273H1	SATMON011	g170772	BLASTN	1364	1e-104	91
1379	503	700573027H1	SATMON030	g170772	BLASTN	939	1e-103	89
1380	503	700209360H1	SATMON016	g170772	BLASTN	1350	1e-103	89
1381	503	700619916H1	SATMON034	g170772	BLASTN	1050	1e-102	89
1382	503	700086051H1	SATMON011	g170772	BLASTN	1336	1e-102	90
1383	503	LIB3069-034-Q1-K1-C8	LIB3069	g170772	BLASTN	1322	1e-101	86
1384	503	700026324H1	SATMON003	g170772	BLASTN	1328	1e-101	91
1385	503	700104549H1	SATMON010	g170772	BLASTN	836	1e-100	88
1386	503	700622108H1	SATMON034	g170772	BLASTN	1158	1e-100	88
1387	503	700093980H1	SATMON008	g170772	BLASTN	1312	1e-100	90
1388	503	700077427H1	SATMON007	g170772	BLASTN	1317	1e-100	90
1389	503	700095389H1	SATMON008	g170772	BLASTN	1302	1e-99	91
1390	503	LIB3067-055-Q1-K1-D3	LIB3067	g170772	BLASTN	762	1e-98	87
1391	503	LIB3060-054-Q1-K1-F6	LIB3060	g170772	BLASTN	1009	1e-98	83
1392	503	700083339H1	SATMON011	g170772	BLASTN	1282	1e-98	91
1393	503	700102631H1	SATMON010	g170772	BLASTN	1283	1e-98	89
1394	503	700265625H1	SATMON017	g170772	BLASTN	1286	1e-98	90
1395	503	700095002H1	SATMON008	g170772	BLASTN	1289	1e-98	88
1396	503	700094761H1	SATMON008	g170772	BLASTN	1289	1e-98	88
1397	503	700073832H1	SATMON007	g170772	BLASTN	1270	1e-97	88
1398	503	700047817H1	SATMON003	g170772	BLASTN	1272	1e-97	91
1399	503	700091149H1	SATMON011	g170772	BLASTN	1262	1e-96	89
1400	503	700098584H1	SATMON009	g170772	BLASTN	1264	1e-96	91
1401	503	700085932H1	SATMON011	g170772	BLASTN	1266	1e-96	89
1402	503	700094081H1	SATMON008	g170772	BLASTN	1269	1e-96	91
1403	503	700202442H1	SATMON003	g170772	BLASTN	1145	1e-95	88
1404	503	700049770H1	SATMON003	g170772	BLASTN	1246	1e-95	89
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1406	503	700090226H1	SATMON011	g170772	BLASTN	1252	1e-95	90
1407	503	700088191H1	SATMON011	g170772	BLASTN	1253	1e-95	90
1408	503	700076743H1	SATMON007	g170772	BLASTN	1256	1e-95	90



1409	503	LIB189-009-Q1-E1-E10	LIB189	g170772	BLASTN	789	1e-94	88
1410	503	700072038H1	SATMON007	g170772	BLASTN	1237	1e-94	89
1411	503	700082967H1	SATMON011	g170772	BLASTN	1239	1e-94	89
1412	503	LIB3068-062-Q1-K1-A3	LIB3068	g170772	BLASTN	1148	1e-93	82
1413	503	700094713H1	SATMON008	g170772	BLASTN	1222	1e-93	88
1414	503	700095620H1	SATMON008	g170772	BLASTN	1173	1e-92	89
1415	503	700242509H1	SATMON010	g170772	BLASTN	1213	1e-92	92
1416	503	700071923H1	SATMON007	g170772	BLASTN	1214	1e-92	90
1417	503	700575314H1	SATMON030	g170772	BLASTN	1149	1e-91	88
1418	503	700086654H1	SATMON011	g170772	BLASTN	1199	1e-91	90
1419	503	700241072H1	SATMON010	g170772	BLASTN	1205	1e-91	91
1420	503	700047361H1	SATMON003	g170772	BLASTN	1110	1e-90	89
1421	503	700217056H1	SATMON016	g170772	BLASTN	1187	1e-90	91
1422	503	LIB3067-005-Q1-K1-F2	LIB3067	g170772	BLASTN	1188	1e-90	84
1423	503	700075495H1	SATMON007	g170772	BLASTN	1193	1e-90	86
1424	503	700084783H1	SATMON011	g170772	BLASTN	1194	1e-90	91
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1427	503	700159079H1	SATMON012	g170772	BLASTN	1177	1e-89	92
1428	503	700094408H1	SATMON008	g170772	BLASTN	1177	1e-89	92
1429	503	700348440H1	SATMON023	g170772	BLASTN	1182	1e-89	89
1430	503	700077082H1	SATMON007	g170772	BLASTN	1182	1e-89	92
1431	503	700082111H1	SATMON011	g170772	BLASTN	1183	1e-89	89
1432	503	700239752H1	SATMON010	g170772	BLASTN	1164	1e-88	89
1433	503	700029456H1	SATMON003	g170772	BLASTN	1166	1e-88	90
1434	503	700209430H1	SATMON016	g170772	BLASTN	1170	1e-88	90
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1436	503	700213138H1	SATMON016	g170772	BLASTN	1042	1e-87	89
1437	503	700102234H1	SATMON010	g170772	BLASTN	634	1e-85	90
1438	503	700450479H1	SATMON028	g170772	BLASTN	1015	1e-85	90
1439	503	700095372H1	SATMON008	g170772	BLASTN	1126	1e-85	90
1440	503	700218734H1	SATMON011	g170772	BLASTN	1128	1e-85	94
1441	503	700256858H1	SATMON017	g170772	BLASTN	1133	1e-85	90
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1443	503	700105439H1	SATMON010	g170772	BLASTN	1118	1e-84	89
1444	503	700085174H1	SATMON011	g170772	BLASTN	1125	1e-84	90
1445	503	700242421H1	SATMON010	g170772	BLASTN	1125	1e-84	92
1446	503	700077492H1	SATMON007	g170772	BLASTN	774	1e-83	88
1447	503	700209295H1	SATMON016	g170772	BLASTN	914	1e-83	89
1448	503	700455826H1	SATMON029	g170772	BLASTN	978	1e-83	91
1449	503	700213370H1	SATMON016	g170772	BLASTN	1110	1e-83	92
1450	503	700352095H1	SATMON023	g170772	BLASTN	1111	1e-83	89
1451	503	700076842H1	SATMON007	g170772	BLASTN	1112	1e-83	90
1452	503	700215872H1	SATMON016	g170772	BLASTN	1113	1e-83	89
1453	503	700073645H1	SATMON007	g170772	BLASTN	1113	1e-83	89
1454	503	700048153H1	SATMON003	g170772	BLASTN	651	1e-82	90
1455	503	700073307H1	SATMON007	g170772	BLASTN	762	1e-82	90
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1458	503	700073955H1	SATMON007	g170772	BLASTN	1090	1e-82	90
1459	503	LIB3059-042-Q1-K1-H12	LIB3059	g170772	BLASTN	1090	1e-82	85
1460	503	700238024H1	SATMON010	g170772	BLASTN	1091	1e-82	90
1461	503	700155863H1	SATMON007	g170772	BLASTN	1091	1e-82	93
1462	503	700217890H1	SATMON016	g170772	BLASTN	1093	1e-82	91

1463	503	700239314H1	SATMON010	g170772	BLASTN	1094	1e-82	88
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1465	503	700235469H1	SATMON010	g170772	BLASTN	1098	1e-82	91
1466	503	700164224H1	SATMON013	g170772	BLASTN	710	1e-81	92
1467	503	700209374H1	SATMON016	g170772	BLASTN	1083	1e-81	89
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1469	503	700159326H1	SATMON012	g170772	BLASTN	777	1e-80	91
1470	503	700025934H1	SATMON003	g170772	BLASTN	792	1e-80	91
1471	503	700210458H1	SATMON016	g170772	BLASTN	856	1e-80	88
1472	503	700243826H1	SATMON010	g170772	BLASTN	881	1e-80	90
1473	503	700242676H1	SATMON010	g170772	BLASTN	940	1e-80	89
1474	503	700345565H1	SATMON021	g170772	BLASTN	1034	1e-80	89
1475	503	700264733H1	SATMON017	g170772	BLASTN	1044	1e-80	90
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1479	503	700451515H1	SATMON028	g170772	BLASTN	1057	1e-79	85
1480	503	700468929H1	SATMON025	g170772	BLASTN	1061	1e-79	84
1481	503	700205801H1	SATMON003	g170772	BLASTN	1064	1e-79	91
1482	503	700611326H1	SATMON022	g170772	BLASTN	725	1e-78	88
1483	503	700082291H1	SATMON011	g170772	BLASTN	1043	1e-78	87
1484	503	700071695H1	SATMON007	g170772	BLASTN	1048	1e-78	89
1485	503	700551561H1	SATMON022	g170772	BLASTN	746	1e-77	88
1486	503	700221019H1	SATMON011	g170772	BLASTN	856	1e-77	89
1487	503	LIB3078-007-Q1-K1-C5	LIB3078	g170772	BLASTN	936	1e-77	83
1488	503	700049925H1	SATMON003	g170772	BLASTN	954	1e-77	89
1489	503	700154101H1	SATMON007	g170772	BLASTN	1036	1e-77	88
1490	503	700380151H1	SATMON021	g170772	BLASTN	613	1e-76	88
1491	503	700243831H1	SATMON010	g170772	BLASTN	842	1e-76	89
1492	503	700235671H1	SATMON010	g170772	BLASTN	1019	1e-76	90
1493	503	700087847H1	SATMON011	g170772	BLASTN	1009	1e-75	82
1494	503	700208747H1	SATMON016	g170772	BLASTN	1010	1e-75	89
1495	503	700071795H1	SATMON007	g170772	BLASTN	1011	1e-75	89
1496	503	700157002H1	SATMON012	g170772	BLASTN	1012	1e-75	90
1497	503	700201608H1	SATMON003	g170772	BLASTN	1015	1e-75	85
1498	503	700451541H1	SATMON028	g170772	BLASTN	1015	1e-75	86
1499	503	700212182H1	SATMON016	g170772	BLASTN	1015	1e-75	88
1500	503	700381485H1	SATMON023	g170772	BLASTN	1017	1e-75	91
1501	503	700096793H1	SATMON008	g170772	BLASTN	761	1e-74	87
1502	503	700093025H1	SATMON008	g170772	BLASTN	996	1e-74	89
1503	503	700017129H1	SATMON001	g170772	BLASTN	998	1e-74	92
1504	503	700216525H1	SATMON016	g170772	BLASTN	1000	1e-74	86
1505	503	700087912H1	SATMON011	g170772	BLASTN	1001	1e-74	91
1506	503	700172618H1	SATMON013	g170772	BLASTN	1001	1e-74	91
1507	503	700801894H1	SATMON036	g170772	BLASTN	1002	1e-74	90
1508	503	700162040H1	SATMON012	g170772	BLASTN	1003	1e-74	92
1509	503	700212084H1	SATMON016	g170772	BLASTN	1004	1e-74	89
1510	503	700027971H1	SATMON003	g170772	BLASTN	982	1e-73	92
1511	503	700077321H1	SATMON007	g170772	BLASTN	985	1e-73	89
1512	503	700619884H1	SATMON034	g170772	BLASTN	987	1e-73	87
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1515	503	700083350H1	SATMON011	g170772	BLASTN	971	1e-72	89
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1517	503	700094688H1	SATMON008	g170772	BLASTN	975	1e-72	89
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1519	503	700451847H1	SATMON028	g170772	BLASTN	872	1e-71	84
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1523	503	700084952H1	SATMON011	g170772	BLASTN	552	1e-70	78
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1530	503	700073939H1	SATMON007	g170772	BLASTN	931	1e-68	88
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1532	503	700072843H1	SATMON007	g170772	BLASTN	692	1e-67	87
1533	503	700439356H1	SATMON026	g170772	BLASTN	777	1e-67	85
1534	503	700073556H1	SATMON007	g170772	BLASTN	910	1e-67	88
1535	503	700154461H1	SATMON007	g170772	BLASTN	911	1e-67	87
1536	503	700201909H1	SATMON003	g170772	BLASTN	912	1e-67	88
1537	503	700104384H1	SATMON010	g170772	BLASTN	665	1e-66	89
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1540	503	700456945H1	SATMON029	g170772	BLASTN	902	1e-66	80
1541	503	700457294H1	SATMON029	g170772	BLASTN	446	1e-65	88
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1543	503	700072428H2	SATMON007	g170772	BLASTN	890	1e-65	88
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1547	503	700240615H1	SATMON010	g170772	BLASTN	877	1e-64	88
1548	503	700155233H1	SATMON007	g170772	BLASTN	883	1e-64	90
1549	503	700241210H1	SATMON010	g170772	BLASTN	822	1e-63	87
1550	503	700478070H1	SATMON025	g170772	BLASTN	863	1e-63	82
1551	503	700447433H1	SATMON027	g170772	BLASTN	865	1e-63	84
1552	503	700215884H1	SATMON016	g170772	BLASTN	865	1e-63	88
1553	503	700575250H1	SATMON030	g170772	BLASTN	820	1e-61	84
1554	503	700209283H1	SATMON016	g170772	BLASTN	833	1e-60	86
1555	503	700213479H1	SATMON016	g170772	BLASTN	834	1e-60	88
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1569	503	700219086H1	SATMON011	g170772	BLASTN	769	1e-55	87
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1571	503	700424527H1	SATMONN01	g170772	BLASTN	777	1e-55	83
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1573	503	700217412H1	SATMON016	g170772	BLASTN	755	1e-54	87
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1576	503	700446645H1	SATMON027	g170772	BLASTN	744	1e-53	87
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1578	503	700617325H1	SATMON033	g170772	BLASTN	751	1e-53	91
1579	503	700239901H1	SATMON010	g170772	BLASTN	641	1e-52	86
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1581	503	700083360H1	SATMON011	g170772	BLASTN	739	1e-52	87
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1583	503	700074780H1	SATMON007	g170772	BLASTN	528	1e-51	83
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1586	503	700165354H1	SATMON013	g170772	BLASTN	728	1e-51	86
1587	503	700353994H1	SATMON024	g170772	BLASTN	713	1e-50	87
1588	503	700153619H1	SATMON007	g170772	BLASTN	694	1e-49	86
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1590	503	700155530H1	SATMON007	g170772	BLASTN	686	1e-48	86
1591	503	700221207H1	SATMON011	g170772	BLASTN	692	1e-48	86
1592	503	700264257H1	SATMON017	g170772	BLASTN	672	1e-47	87
1593	503	700152216H1	SATMON007	g170772	BLASTN	659	1e-46	88
1594	503	700150643H1	SATMON007	g170772	BLASTN	659	1e-46	88
1595	503	700156027H1	SATMON007	g170772	BLASTN	655	1e-45	93
1596	503	700260335H1	SATMON017	g170772	BLASTN	385	1e-44	79
1597	503	700623795H1	SATMON034	g170772	BLASTN	401	1e-44	88
1598	503	700150581H1	SATMON007	g170772	BLASTN	640	1e-44	88
1599	503	700151970H1	SATMON007	g170772	BLASTN	645	1e-44	87
1600	503	700151780H1	SATMON007	g170772	BLASTN	628	1e-43	88
1601	503	700575547H1	SATMON030	g170772	BLASTN	614	1e-42	90
1602	503	LIB143-026-Q1-E1-A5	LIB143	g170772	BLASTN	633	1e-42	77
1603	503	700347945H1	SATMON023	g170772	BLASTN	586	1e-40	84
1604	503	700074416H1	SATMON007	g170772	BLASTN	589	1e-40	86
1605	503	700352990H1	SATMON024	g170772	BLASTN	575	1e-39	92
1606	503	700156025H1	SATMON007	g170772	BLASTN	568	1e-38	91
1607	503	700432072H1	SATMONN01	g170772	BLASTN	374	1e-37	84
1608	503	700354249H1	SATMON024	g170772	BLASTN	528	1e-35	93
1609	503	700617977H1	SATMON033	g170772	BLASTN	535	1e-35	87
1610	503	700218312H1	SATMON016	g170772	BLASTN	236	1e-33	91
1611	503	700456080H1	SATMON029	g170772	BLASTN	513	1e-33	83
1612	503	700349395H1	SATMON023	g170772	BLASTN	500	1e-32	91
1613	503	700051637H1	SATMON003	g170772	BLASTN	249	1e-31	86
1614	503	700202153H1	SATMON003	g170772	BLASTN	474	1e-30	91
1615	503	700256951H1	SATMON017	g170772	BLASTN	454	1e-29	86
1616	503	700150542H1	SATMON007	g170772	BLASTN	343	1e-28	76
1617	503	700151629H1	SATMON007	g170772	BLASTN	445	1e-28	85
1618	503	700446654H1	SATMON027	g170772	BLASTN	437	1e-27	84
1619	503	700155814H1	SATMON007	g170772	BLASTN	428	1e-26	88
1620	503	700161019H1	SATMON012	g2588780	BLASTN	417	1e-25	92
1621	503	700377236H1	SATMON019	g170772	BLASTN	402	1e-24	84
1622	503	LIB3067-036-Q1-K1-A4	LIB3067	g1220121	BLASTN	224	1e-22	84
1623	503	700158933H1	SATMON012	g170772	BLASTN	368	1e-21	90
1624	503	700155365H1	SATMON007	g170772	BLASTN	347	1e-20	91

1625	503	700405366H1	SATMON029	g170772	BLASTN	311	1e-17	88
1626	503	700159812H1	SATMON012	g407412	BLASTX	160	1e-15	96
1627	503	700209759H1	SATMON016	g170772	BLASTN	239	1e-15	90
1628	503	700154527H1	SATMON007	g170772	BLASTN	250	1e-12	92
1629	503	700449637H1	SATMON028	g170772	BLASTN	201	1e-10	78
1630	503	700096829H1	SATMON008	g170772	BLASTN	216	1e-9	89
2993	-700661285	700661285H1	SOYMON005	g1857024	BLASTX	95	1e-12	100
2994	-700750570	700750570H1	SOYMON014	g170772	BLASTN	414	1e-24	81
2995	-700752735	700752735H1	SOYMON014	g170772	BLASTN	446	1e-27	78
2996	-700755052	700755052H1	SOYMON014	g170772	BLASTN	547	1e-45	74
2997	-700756501	700756501H1	SOYMON014	g535583	BLASTN	717	1e-50	84
2998	-700831127	700831127H1	SOYMON019	g535583	BLASTN	862	1e-63	83
2999	-700851779	700851779H1	SOYMON023	g170772	BLASTN	505	1e-33	77
3000	-700888715	700888715H1	SOYMON024	g535583	BLASTN	442	1e-31	91
3001	-700889420	700889420H1	SOYMON024	g1220121	BLASTN	893	1e-65	84
3002	-700895218	700895218H1	SOYMON024	g407411	BLASTN	816	1e-59	83
3003	-700941379	700941379H1	SOYMON024	g170772	BLASTN	424	1e-33	72
3004	-700986855	700986855H1	SOYMON009	g170772	BLASTN	701	1e-49	74
3005	-701070484	701070484H1	SOYMON034	g407411	BLASTN	362	1e-43	73
3006	-701136279	701136279H1	SOYMON038	g170772	BLASTN	651	1e-56	80
3007	-GM16478	LIB3054-007-Q1-N1-G3	LIB3054	g2244750	BLASTX	70	1e-27	61
3008	-GM23819	LIB3040-019-Q1-E1-C5	LIB3040	g535583	BLASTN	258	1e-10	88
3009	-GM29758	LIB3050-016-Q1-E1-B11	LIB3050	g535583	BLASTN	391	1e-42	70
3010	16	LIB3030-003-Q1-B1-B11	LIB3030	g3088578	BLASTN	1577	1e-122	87
3011	16	LIB3050-023-Q1-K1-H9	LIB3050	g535583	BLASTN	1498	1e-116	86
3012	16	LIB3030-003-Q1-B1-F7	LIB3030	g535583	BLASTN	1469	1e-113	86
3013	16	LIB3055-005-Q1-N1-C11	LIB3055	g170772	BLASTN	1205	1e-109	84
3014	16	LIB3065-011-Q1-N1-A3	LIB3065	g170772	BLASTN	622	1e-107	88
3015	16	700652256H1	SOYMON003	g170772	BLASTN	591	1e-90	88
3016	16	LIB3065-011-Q1-N1-A4	LIB3065	g170772	BLASTN	1182	1e-89	78
3017	16	700653827H1	SOYMON003	g170772	BLASTN	660	1e-87	81
3018	16	701099940H1	SOYMON028	g1220121	BLASTN	1154	1e-87	88
3019	16	701003671H1	SOYMON019	g170772	BLASTN	1131	1e-85	92
3020	16	700752105H1	SOYMON014	g170772	BLASTN	1116	1e-84	91
3021	16	700653057H1	SOYMON003	g170772	BLASTN	990	1e-83	88
3022	16	700945531H1	SOYMON024	g535583	BLASTN	1109	1e-83	89
3023	16	700980013H1	SOYMON009	g535583	BLASTN	1109	1e-83	88
3024	16	701127812H1	SOYMON037	g170772	BLASTN	1110	1e-83	91
3025	16	LIB3056-014-Q1-N1-F8	LIB3056	g170772	BLASTN	798	1e-81	82
3026	16	700653862H1	SOYMON003	g1220121	BLASTN	960	1e-80	88
3027	16	700994148H1	SOYMON011	g535583	BLASTN	1069	1e-80	86
3028	16	700984184H1	SOYMON009	g170772	BLASTN	756	1e-79	86
3029	16	701123715H1	SOYMON037	g1220121	BLASTN	1059	1e-79	87
3030	16	700839038H1	SOYMON020	g170772	BLASTN	1060	1e-79	93
3031	16	700978445H1	SOYMON009	g170772	BLASTN	1062	1e-79	89
3032	16	701123035H1	SOYMON037	g170772	BLASTN	829	1e-78	90
3033	16	701041545H1	SOYMON029	g170772	BLASTN	1030	1e-77	86
3034	16	700898192H1	SOYMON027	g1220121	BLASTN	1031	1e-77	89
3035	16	700985750H1	SOYMON009	g169662	BLASTN	1033	1e-77	85
3036	16	700730995H1	SOYMON009	g170772	BLASTN	1040	1e-77	91
3037	16	700555909H1	SOYMON001	g535583	BLASTN	593	1e-76	84
3038	16	700746538H1	SOYMON013	g535583	BLASTN	834	1e-76	88
3039	16	700941380H1	SOYMON024	g170772	BLASTN	991	1e-76	91
3040	16	701118851H1	SOYMON037	g170772	BLASTN	1019	1e-76	89

3041	16	701065379H1	SOYMON034	g535583	BLASTN	1022	1e-76	86
3042	16	701209645H1	SOYMON035	g170772	BLASTN	795	1e-75	86
3043	16	701055017H1	SOYMON032	g1220121	BLASTN	877	1e-75	86
3044	16	701015213H1	SOYMON019	g535583	BLASTN	1008	1e-75	87
3045	16	700982770H1	SOYMON009	g1220121	BLASTN	1009	1e-75	85
3046	16	701212420H1	SOYMON035	g535583	BLASTN	1012	1e-75	86
3047	16	700977916H1	SOYMON009	g170772	BLASTN	700	1e-74	89
3048	16	700974401H1	SOYMON005	g1220121	BLASTN	857	1e-74	89
3049	16	700645749H1	SOYMON010	g170772	BLASTN	996	1e-74	82
3050	16	700646620H1	SOYMON014	g170772	BLASTN	996	1e-74	89
3051	16	701126103H1	SOYMON037	g170772	BLASTN	1000	1e-74	88
3052	16	700978001H1	SOYMON009	g170772	BLASTN	563	1e-73	89
3053	16	700561987H1	SOYMON002	g1220121	BLASTN	782	1e-73	87
3054	16	701101526H1	SOYMON028	g1220121	BLASTN	810	1e-73	88
3055	16	700562680H1	SOYMON002	g170772	BLASTN	985	1e-73	87
3056	16	700560521H1	SOYMON001	g170772	BLASTN	988	1e-73	80
3057	16	701055544H1	SOYMON032	g170772	BLASTN	992	1e-73	88
3058	16	701061418H1	SOYMON033	g170772	BLASTN	993	1e-73	87
3059	16	701049514H1	SOYMON032	g1220121	BLASTN	884	1e-72	88
3060	16	700646215H1	SOYMON012	g170772	BLASTN	971	1e-72	89
3061	16	701014875H1	SOYMON019	g535583	BLASTN	973	1e-72	87
3062	16	700874718H1	SOYMON018	g169662	BLASTN	974	1e-72	86
3063	16	700897796H1	SOYMON027	g1220121	BLASTN	977	1e-72	87
3064	16	700548019H1	SOYMON001	g170772	BLASTN	547	1e-71	87
3065	16	700904875H1	SOYMON022	g535583	BLASTN	963	1e-71	88
3066	16	701106315H1	SOYMON036	g170772	BLASTN	557	1e-70	90
3067	16	700745023H1	SOYMON013	g407411	BLASTN	949	1e-70	85
3068	16	701120348H1	SOYMON037	g170772	BLASTN	956	1e-70	86
3069	16	701124508H1	SOYMON037	g170772	BLASTN	578	1e-69	84
3070	16	700956183H1	SOYMON022	g1220121	BLASTN	625	1e-69	87
3071	16	700894619H1	SOYMON024	g535583	BLASTN	828	1e-69	85
3072	16	700892392H1	SOYMON024	g535583	BLASTN	936	1e-69	85
3073	16	700730676H1	SOYMON009	g535583	BLASTN	939	1e-69	86
3074	16	700728913H1	SOYMON009	g407411	BLASTN	940	1e-69	87
3075	16	700845737H1	SOYMON021	g1220121	BLASTN	528	1e-68	88
3076	16	700990961H1	SOYMON011	g170772	BLASTN	582	1e-68	85
3077	16	700983443H1	SOYMON009	g170772	BLASTN	922	1e-68	86
3078	16	701120754H1	SOYMON037	g170772	BLASTN	923	1e-68	88
3079	16	700900486H1	SOYMON027	g1220121	BLASTN	924	1e-68	83
3080	16	701133574H2	SOYMON038	g170772	BLASTN	929	1e-68	88
3081	16	700900924H1	SOYMON027	g170772	BLASTN	931	1e-68	88
3082	16	701056706H1	SOYMON032	g170772	BLASTN	486	1e-67	89
3083	16	701110051H1	SOYMON036	g170772	BLASTN	910	1e-67	88
3084	16	701136325H1	SOYMON038	g170772	BLASTN	915	1e-67	87
3085	16	700750809H1	SOYMON014	g170772	BLASTN	900	1e-66	88
3086	16	700686607H1	SOYMON008	g170772	BLASTN	901	1e-66	88
3087	16	700848261H1	SOYMON021	g535583	BLASTN	905	1e-66	87
3088	16	700686634H1	SOYMON008	g170772	BLASTN	905	1e-66	88
3089	16	700891285H1	SOYMON024	g170772	BLASTN	906	1e-66	88
3090	16	700560291H1	SOYMON001	g170772	BLASTN	906	1e-66	88
3091	16	700752975H1	SOYMON014	g170772	BLASTN	907	1e-66	89
3092	16	701006013H2	SOYMON019	g170772	BLASTN	908	1e-66	87
3093	16	700974038H1	SOYMON005	g1220121	BLASTN	631	1e-65	88
3094	16	701047024H1	SOYMON032	g170772	BLASTN	707	1e-65	89

3095	16	700900409H1	SOYMON027	g170772	BLASTN	737	1e-65	83
3096	16	701137320H1	SOYMON038	g170772	BLASTN	769	1e-65	88
3097	16	700978805H1	SOYMON009	g170772	BLASTN	886	1e-65	84
3098	16	700726195H1	SOYMON009	g1220121	BLASTN	889	1e-65	87
3099	16	700661112H1	SOYMON005	g170772	BLASTN	661	1e-64	85
3100	16	700989712H1	SOYMON011	g170772	BLASTN	788	1e-64	88
3101	16	700752287H1	SOYMON014	g170772	BLASTN	875	1e-64	85
3102	16	700964226H1	SOYMON022	g170772	BLASTN	877	1e-64	88
3103	16	700847346H1	SOYMON021	g170772	BLASTN	880	1e-64	83
3104	16	700756428H1	SOYMON014	g170772	BLASTN	883	1e-64	88
3105	16	701049928H1	SOYMON032	g170772	BLASTN	885	1e-64	82
3106	16	700848652H1	SOYMON021	g170772	BLASTN	477	1e-63	89
3107	16	700898929H1	SOYMON027	g1220121	BLASTN	514	1e-63	87
3108	16	700903523H1	SOYMON022	g170772	BLASTN	527	1e-63	82
3109	16	700983745H1	SOYMON009	g1220121	BLASTN	863	1e-63	88
3110	16	700890587H1	SOYMON024	g170772	BLASTN	865	1e-63	86
3111	16	700969917H1	SOYMON005	g407411	BLASTN	868	1e-63	83
3112	16	700808487H1	SOYMON024	g170772	BLASTN	870	1e-63	88
3113	16	700749968H1	SOYMON013	g170772	BLASTN	871	1e-63	87
3114	16	700751254H1	SOYMON014	g170772	BLASTN	575	1e-62	87
3115	16	701014277H1	SOYMON019	g170772	BLASTN	739	1e-62	86
3116	16	700853635H1	SOYMON023	g170772	BLASTN	854	1e-62	87
3117	16	700752357H1	SOYMON014	g170772	BLASTN	856	1e-62	88
3118	16	700754523H1	SOYMON014	g170772	BLASTN	856	1e-62	92
3119	16	700982153H1	SOYMON009	g170772	BLASTN	400	1e-61	82
3120	16	700958283H1	SOYMON022	g170772	BLASTN	839	1e-61	87
3121	16	700980911H1	SOYMON009	g170772	BLASTN	842	1e-61	82
3122	16	700788112H1	SOYMON011	g170772	BLASTN	844	1e-61	83
3123	16	701005927H1	SOYMON019	g170772	BLASTN	849	1e-61	88
3124	16	700756443H1	SOYMON014	g170772	BLASTN	849	1e-61	88
3125	16	700658914H1	SOYMON004	g1220121	BLASTN	468	1e-60	85
3126	16	701135266H1	SOYMON038	g170772	BLASTN	827	1e-60	87
3127	16	700982179H1	SOYMON009	g170772	BLASTN	827	1e-60	82
3128	16	700754593H1	SOYMON014	g170772	BLASTN	832	1e-60	81
3129	16	700831723H1	SOYMON019	g170772	BLASTN	814	1e-59	91
3130	16	700986775H1	SOYMON009	g170772	BLASTN	815	1e-59	92
3131	16	700755219H1	SOYMON014	g170772	BLASTN	820	1e-59	84
3132	16	701015494H1	SOYMON019	g170772	BLASTN	822	1e-59	88
3133	16	701008473H1	SOYMON019	g170772	BLASTN	823	1e-59	87
3134	16	700754981H1	SOYMON014	g170772	BLASTN	824	1e-59	88
3135	16	700870790H1	SOYMON018	g170772	BLASTN	825	1e-59	81
3136	16	700833069H1	SOYMON019	g170772	BLASTN	806	1e-58	86
3137	16	700680127H2	SOYMON008	g535583	BLASTN	807	1e-58	86
3138	16	701015374H1	SOYMON019	g170772	BLASTN	807	1e-58	87
3139	16	700872895H1	SOYMON018	g535583	BLASTN	808	1e-58	88
3140	16	701137912H1	SOYMON038	g170772	BLASTN	374	1e-57	83
3141	16	700984063H1	SOYMON009	g1220121	BLASTN	575	1e-57	79
3142	16	700991988H1	SOYMON011	g170772	BLASTN	791	1e-57	82
3143	16	700873915H1	SOYMON018	g170772	BLASTN	793	1e-57	88
3144	16	700978721H1	SOYMON009	g170772	BLASTN	797	1e-57	78
3145	16	701213679H1	SOYMON035	g170772	BLASTN	798	1e-57	88
3146	16	701102954H1	SOYMON028	g170772	BLASTN	799	1e-57	79
3147	16	700888289H1	SOYMON024	g170772	BLASTN	712	1e-56	91
3148	16	700962086H1	SOYMON022	g170772	BLASTN	783	1e-56	88

3149	16	701052227H1	SOYMON032	g535583	BLASTN	788	1e-56	86
3150	16	700755177H1	SOYMON014	g170772	BLASTN	768	1e-55	88
3151	16	701123136H1	SOYMON037	g170772	BLASTN	771	1e-55	82
3152	16	700979067H1	SOYMON009	g170772	BLASTN	776	1e-55	90
3153	16	700755513H1	SOYMON014	g170772	BLASTN	759	1e-54	92
3154	16	700756639H1	SOYMON014	g170772	BLASTN	761	1e-54	81
3155	16	700554077H1	SOYMON001	g170772	BLASTN	371	1e-53	86
3156	16	700653194H1	SOYMON003	g170772	BLASTN	395	1e-53	86
3157	16	701110348H1	SOYMON036	g170772	BLASTN	743	1e-53	81
3158	16	700753973H1	SOYMON014	g170772	BLASTN	743	1e-53	87
3159	16	701011009H1	SOYMON019	g535583	BLASTN	733	1e-52	81
3160	16	700739086H1	SOYMON012	g170772	BLASTN	456	1e-51	87
3161	16	700740140H1	SOYMON012	g170772	BLASTN	723	1e-51	89
3162	16	700565040H1	SOYMON002	g170772	BLASTN	729	1e-51	74
3163	16	701148186H1	SOYMON031	g535583	BLASTN	665	1e-50	85
3164	16	701142734H1	SOYMON038	g535583	BLASTN	670	1e-47	86
3165	16	700754441H1	SOYMON014	g170772	BLASTN	385	1e-46	93
3166	16	701102588H1	SOYMON028	g1220121	BLASTN	662	1e-46	89
3167	16	700900656H1	SOYMON027	g535583	BLASTN	635	1e-44	87
3168	16	700974207H1	SOYMON005	g535583	BLASTN	641	1e-44	85
3169	16	700982081H1	SOYMON009	g170772	BLASTN	494	1e-43	78
3170	16	701130026H1	SOYMON037	g1220121	BLASTN	482	1e-42	86
3171	16	701009720H1	SOYMON019	g170772	BLASTN	621	1e-42	87
3172	16	700962602H1	SOYMON022	g170772	BLASTN	357	1e-40	91
3173	16	700724934H1	SOYMON009	g535583	BLASTN	561	1e-40	81
3174	16	700729305H1	SOYMON009	g535583	BLASTN	544	1e-39	82
3175	16	701210054H1	SOYMON035	g535583	BLASTN	569	1e-38	85
3176	16	700790192H1	SOYMON011	g535583	BLASTN	293	1e-37	83
3177	16	700984076H1	SOYMON009	g170772	BLASTN	198	1e-35	88
3178	16	700726562H1	SOYMON009	g170772	BLASTN	307	1e-34	78
3179	16	701211376H1	SOYMON035	g170772	BLASTN	524	1e-34	86
3180	16	700753085H1	SOYMON014	g2588780	BLASTN	358	1e-33	76
3181	16	700727993H1	SOYMON009	g535583	BLASTN	465	1e-32	84
3182	16	700561072H1	SOYMON001	g170772	BLASTN	473	1e-30	83
3183	16	701211464H1	SOYMON035	g170772	BLASTN	464	1e-28	81
3184	16	701098045H1	SOYMON028	g169660	BLASTN	386	1e-21	78
3185	16	700945233H1	SOYMON024	g407412	BLASTX	150	1e-18	87
3186	16	700752655H1	SOYMON014	g758247	BLASTX	172	1e-16	94
3187	16	700735356H1	SOYMON010	g758247	BLASTX	152	1e-14	100
3188	16	700683995H1	SOYMON008	g758247	BLASTX	106	1e-13	90
3189	16	700658760H1	SOYMON004	g1857024	BLASTX	123	1e-13	63
3190	16	700762885H1	SOYMON015	g1857024	BLASTX	134	1e-13	89
3191	16	700755740H1	SOYMON014	g170773	BLASTX	149	1e-13	100
3192	16	700854969H1	SOYMON023	g170772	BLASTN	178	1e-12	80
3193	16	701143036H1	SOYMON038	g169661	BLASTX	113	1e-8	83
3194	18409	700786561H1	SOYMON011	g535583	BLASTN	992	1e-73	85
3195	18409	701008057H1	SOYMON019	g535583	BLASTN	831	1e-60	87
3196	18409	701037442H1	SOYMON029	g535583	BLASTN	669	1e-46	86
3197	18409	700942865H1	SOYMON024	g535583	BLASTN	464	1e-32	86
3198	7322	700651524H1	SOYMON003	g170772	BLASTN	466	1e-65	80
3199	7322	700565758H1	SOYMON002	g170772	BLASTN	450	1e-63	81



**CYSTATHIONINE  $\beta$ -SYNTHASE ( EC 4.2.1.22 )**

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1631	-700025795	700025795H1	SATMON003	g1323263	BLASTX	186	1e-25	68
1632	20651	700344783H1	SATMON021	g1813975	BLASTX	41	1e-9	53

**CYSTATHIONINE  $\gamma$ -LYASE ( EC 4.4.1.1 )**

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
1633	-700260027	700260027H1	SATMON017	g169475	BLASTX	112	1e-10	75
1634	1228	700027629H1	SATMON003	g169475	BLASTX	189	1e-19	87
3203	-700750583	700750583H1	SOYMON014	g169475	BLASTX	149	1e-13	78
3204	12502	LIB3051-069-Q1-K1-E6	LIB3051	g2641242	BLASTX	86	1e-30	38

**O-ACETYLHOMOSERINE (THIOL)-LYASE ( EC 4.2.99.10 )**

Seq No.	Cluster ID	CloneID	Library	NCBI gi	Method	Score	P-value	%Ident
3200	12502	701135185H1	SOYMON038	g1628606	BLASTX	100	1e-10	48
3201	12502	701042913H1	SOYMON029	g2605905	BLASTX	110	1e-9	42
3202	12502	701059330H1	SOYMON033	g2605905	BLASTX	110	1e-9	42

### **\*Table Headings**

#### **Cluster ID**

A cluster ID is arbitrarily assigned to all of those clones which belong to the same cluster at a given stringency and a particular clone will belong to only one cluster at a given stringency. If a cluster contains only a single clone (a “singleton”), then the cluster ID number will be negative, with an absolute value equal to the clone ID number of its single member. The cluster ID entries in the table refer to the cluster with which the particular clone in each row is associated.

#### **Clone ID**

The clone ID number refers to the particular clone in the PhytoSeq database. Each clone ID entry in the table refers to the clone whose sequence is used for (1) the sequence comparison whose scores are presented and/or (2) assignment to the particular cluster which is presented. Note that a clone may be included in this table even if its sequence comparison scores fail to meet the minimum standards for similarity. In such a case, the clone is included due solely to its association with a particular cluster for which sequences of one or more other member clones possess the required level of similarity.

#### **Library**

The library ID refers to the particular cDNA library from which a given clone is obtained. Each cDNA library is associated with the particular tissue(s), line(s) and developmental stage(s) from which it is isolated.

#### **NCBI gi**

Each sequence in the GenBank public database is arbitrarily assigned a unique NCBI gi (National Center for Biotechnology Information GenBank Identifier) number. In this table, the

NCBI gi number which is associated (in the same row) with a given clone refers to the particular GenBank sequence which is used in the sequence comparison. This entry is omitted when a clone is included solely due to its association with a particular cluster.

### **Method**

The entry in the “Method” column of the table refers to the type of BLAST search that is used for the sequence comparison. “CLUSTER” is entered when the sequence comparison scores for a given clone fail to meet the minimum values required for significant similarity. In such cases, the clone is listed in the table solely as a result of its association with a given cluster for which sequences of one or more other member clones possess the required level of similarity.

### **Score**

Each entry in the “Score” column of the table refers to the BLAST score that is generated by sequence comparison of the designated clone with the designated GenBank sequence using the designated BLAST method. This entry is omitted when a clone is included solely due to its association with a particular cluster. If the program used to determine the hit is HMMSW then the score refers to HMMSW score.

### **P-Value**

The entries in the P-Value column refer to the probability that such matches occur by chance.

### **%Ident**

The entries in the “%Ident” column of the table refer to the percentage of identically matched nucleotides (or residues) that exist along the length of that portion of the sequences which is aligned by the BLAST comparison to generate the statistical scores presented. This entry is omitted when a clone is included solely due to its association with a particular cluster.

**We claim:**

1. A substantially purified nucleic acid molecule that encodes a maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of:

- (a) methionine adenosyltransferase,
- (b) S-adenosyl-methionine decarboxylase,
- (c) aspartate kinase,
- (d) aspartate-semialdehyde dehydrogenase,
- (e) cystathionine gamma-synthase,
- (f) cystathionine beta-lysase, and
- (g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase.

2. The substantially purified nucleic acid molecule according to claim 1, wherein said nucleic acid molecule comprises a nucleic acid sequence selected from the group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204.

3. A substantially purified maize or soybean enzyme or fragment thereof, wherein said maize or soybean enzyme is selected from the group consisting of

- (a) methionine adenosyltransferase or fragment thereof;
- (b) S-adenosyl-methionine decarboxylase or fragment thereof;
- (c) aspartate kinase or fragment thereof;
- (d) aspartate-semialdehyde dehydrogenase or fragment thereof;
- (e) cystathionine gamma-synthase or fragment thereof;
- (f) cystathionine beta-lysase or fragment thereof; and

(g) 5-methyltetrahydropteroyltriglutamate-homocysteine-S-methyltransferase or fragment thereof.

4. A substantially purified maize or soybean enzyme or fragment thereof according to claim 3, wherein said maize or soybean enzyme or fragment thereof is encoded by a nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of consisting of SEQ ID NO: 1 through SEQ ID NO: 3204.

5. A substantially purified antibody or fragment thereof which is capable of specifically binding to a specific maize or soybean enzyme or fragment thereof according to claim 4.

6. A transformed plant having a nucleic acid molecule which comprises:

(A) an exogenous promoter region which functions in a plant cell to cause the production of a mRNA molecule;

(B) a structural nucleic acid molecule comprising a nucleic acid sequence selected from the group consisting of

(a) a nucleic acid sequence which encodes for methionine adenosyltransferase or fragment thereof;

(b) a nucleic acid sequence which encodes for S-adenosyl-methionine decarboxylase or fragment thereof;

(c) a nucleic acid sequence which encodes for aspartate kinase or fragment thereof;

(d) a nucleic acid sequence which encodes for aspartate-semialdehyde dehydrogenase or fragment thereof;

(e) a nucleic acid sequence which encodes for cystathionine gamma-synthase or a fragment thereof;

(f) a nucleic acid sequence which encodes for cystathionine beta-lyase or a fragment thereof;

(g) a nucleic acid sequence which encodes for 5-methyltetrahydropteroyl-triglutamate-homocysteine-S-methyltransferase or a fragment thereof; and

(h) a nucleic acid sequence which is complementary to any of the nucleic acid sequences of (a) through (g); and

(C) a 3' non-translated sequence that functions in said plant cell to cause termination of transcription and addition of polyadenylated ribonucleotides to a 3' end of said mRNA molecule.

7. The transformed plant according to claim 5, wherein said structural gene is complementary to any of the nucleic acid sequences of (a) through (g).

8. A method for determining a level or pattern in a plant cell of an enzyme in a plant metabolic pathway comprising:

(A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid molecule, said marker nucleic acid molecule selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having the nucleic acid sequence of SEQ ID NO: 1 through SEQ ID NO: 3204 or compliments thereof, with a complementary nucleic acid molecule obtained from said plant cell or plant tissue, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue permits the detection of an mRNA for said enzyme;

(B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant cell or plant tissue; and

(C) detecting the level or pattern of said complementary nucleic acid, wherein the detection of said complementary nucleic acid is predictive of the level or pattern of said enzyme in said plant metabolic pathway.

9. The method of claim 8, wherein said level or pattern is detected by *in situ* hybridization.

10. A method of determining a mutation in a plant whose presence is predictive of a mutation affecting a level or pattern of a protein comprising the steps:

(A) incubating, under conditions permitting nucleic acid hybridization, a marker nucleic acid, said marker nucleic acid selected from the group of marker nucleic acid molecules which specifically hybridize to a nucleic acid molecule having a nucleic acid sequence selected from the group of SEQ ID NO: 1 through SEQ ID NO: 3204 or complements thereof and a complementary nucleic acid molecule obtained from said plant, wherein nucleic acid hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant permits the detection of a polymorphism whose presence is predictive of a mutation affecting said level or pattern of said plant methionine pathway enzyme in said plant;

(B) permitting hybridization between said marker nucleic acid molecule and said complementary nucleic acid molecule obtained from said plant; and

(C) detecting the presence of said polymorphism, wherein the detection of said polymorphism is predictive of said mutation.

11. A method of producing a plant containing an overexpressed protein comprising:

(A) transforming said plant with a functional nucleic acid molecule, wherein the functional nucleic acid molecule comprises a promoter region, wherein said promoter region is linked to a structural region, wherein said structural region has a nucleic acid sequence selected from group consisting of SEQ ID NO: 1 through SEQ ID NO: 3204 wherein said structural region is linked to a 3' non-translated sequence that functions in the plant to cause termination of





## **ABSTRACT**

The present invention is in the field of plant biochemistry. More specifically the invention relates to nucleic acid sequences from plant cells, in particular, DNA sequences from maize and soybean plants associated with the methionine pathway. The invention encompasses nucleic acid molecules that encode proteins and fragments of proteins. In addition, the invention also encompasses proteins and fragments of proteins so encoded and antibodies capable of binding these proteins or fragments. The invention also relates to methods of using the nucleic acid molecules, proteins and fragments of proteins and antibodies, for example for genome mapping, gene identification and analysis, plant breeding, preparation of constructs for use in plant gene expression and transgenic plants.

<110> Bledig, Stefan A.  
 Byrum, Joseph R.  
 Liu, Jingdong  
 Hinkle, Gregory J.

<120> Nucleic Acid molecules And Other Molecules Associated With The  
 Methionine Synthesis And Degradation Pathways

<130> 38-21(15077)B

<160> 3204

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 tnnnnnnnnnn nnnnnnnnnnn nnnnnnnnnnn nnnnnnnnca gggtcatcaa gaccgccgca 180  
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 tgnannaggc gcancccgac aagctgtgcg accaggtgtc ggacgccgtg nttgacgcnt 180  
 gcctcncnca gnaccccgac ancaangtng cctgncacan gtgcaccaag accaacntgg 240  
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cgatgacacg gactattctt ggagagcaaa gggcttccat gagcaagcaa aagctgccgg 180  
tggtccggct attactactg ctggcatcta tccaggagtt agcaatgtga tggctgctga 240  
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<211> 211

<212> nucleic acid

<213> Zea mays

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gttcgtnaac tcgtacgagc accggcacgn tnaangncaa ggagatcntc aanatcntna 180  
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<211> 266

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<213> Zea mays

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ctgacaagat cgctgatcag gtctctgacg ccgttctcga tgctgtcttg gccgaggacc 180

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 cttggcaatg attaataaat caaacctttg aacctccaaa gaaaaanggc gggaaacngg 420  
 cggctaacct cnaanaacgg gcaagcctaa tggggcaact tcggaaagg gaaaccctgg 480  
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tggcgccttc	tcnngnnng	gnccctagca	ncggtngacc	gcancgggcg	ctantgnccc	420
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<212>	nucleic acid
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ctcaccggcc	ancaagatca	tcctngacac	ctacggcnn	tggggagccc	acnncgnggg	180
cgccttcttc	ggnaaggacc	ccaccaaggt	gnaccnnaa	ggggcctacn	tcgccnnnca	240
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cgncatcggc	gtgcctgaan	cnccttcant	nttcgttgaa	ttctanagna	accgggacca	360
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<210>	12
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 ggtgcttggt aacatttaca ncaatccccc acattcccaa ggcnttcacn ggcactttaa 180  
 caaaacgncc naaggaaaat cggccccgcn aacaaagcca aatttttcgg ttccccaccc 240  
 ancanacncc caacttattc ccttaacca cctgcttgca ccaaactttt tnnccccct 300  
 tantcgaagt tcnnaaagga cngnaccttt cctttgttaa accccaaagn aaaaaccaa 360  
 tttaccgtn nantntntta aaaaaaggcc gncacctntt ncccccttc cnttcacaac 420  
 cctnttttat tttttannca aaaa 444

G  
 e  
 n  
 e  
 b  
 a  
 n  
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 G  
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 e  
 b  
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 n  
 k

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 <212> nucleic acid  
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<400> 14

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aaccggtccg ggcgtttcgt catcggcggg cccacgggtg acgccggcct caccggccgc 180  
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ccggagcccc tgtccgtgtt cgtcaactcg tacggcaccg gcacgatccc cgacaaggag 420  
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<211> 511

<212> nucleic acid

<213> Zea mays

<400> 15

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cacagtctc atctctaccc agcacgacga gacagtcacc aacgacgaga ttgctgctga 300  
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<211> 483

<212> nucleic acid

<213> Zea mays

<400> 16

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<210>  
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17  
 454  
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 Zea mays

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18  
 474  
 nucleic acid  
 Zea mays

<400> 18

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 accgacnaga cccccagct gatgccgctc agccacgtgc tggccaccaa gctgggcgcg 300  
 cgcctcaccg aggtgcncan gaacggcacc tgcncctggc tgaggcccga cggcaagacc 360  
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<210> 19  
 <211> 435  
 <212> nucleic acid  
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<400> 19  
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 aaccgtccg ggcgcttcgt catcggcggg cccacgggtg acgcccgcct caccggccgc 300

aagatcatca tgcacacgta cggcggctgg ggaacccaag gcggtggcgc cttctccggc 360  
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<212> nucleic acid  
<213> Zea mays  
<400> 21

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ggcctcgacg ccgaccgctg caaggtgctg gtgaacatcg agcagcaatc cccgacatcg 420  
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<210> 22  
<211> 491  
<212> nucleic acid  
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<400> 22

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accttgacga gaagaccatc ttccacctta acccatctgg ccgctttgtc attggtggac 180  
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cgcagcggag cctatgtcgc aaggcaggct gccaaagagca tcgtcgccag cggccttgct 180  
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ganggggggg aancctta 439

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<211> 502  
<212> nucleic acid  
<213> Zea mays  
<400> 24

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 <213> Zea mays  
  
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<210> 26  
 <211> 393  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 26

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 ctcagggtgt gcatggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 240  
 gacacatgtt cgggtatgcg accgatgaga ccctgagtt gatgcccctc agccatgtcc 300  
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 tcaagcctga tgggaaaaac caaggtgaca gtc 393

<210> 27  
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tgaccgcagc ggagcctatg tcgcaaggca ggctgccaaag agcatcgtnn ccagcggcct 300  
tgctcgccgc gccatcgctc aggtgtctta cgccatcggn gtgcccagac ctctctccgn 360  
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agaactttta tttca 435

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<400> 28

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gaccgcgcga taaggncact ttggccgtga cgacgccgac ttcacctggg aggtggtcaa 360  
gcccctaaag aangcattcc gcttaagaat gtattgggaa gttcactgga catgaagtcc 420  
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<212> nucleic acid  
<213> Zea mays  
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agaccgtcac caacgacgag atcgccgccg acctcaagga gcacgtcatc aagcccgtga 240  
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 tcatcgggcg gccccacggt gacgccggcc tcaccggccg caagatnata atcgacacgt 360  
 acggcggttg gggagccac ggcggtggcg ccttctccgg caaggacccc accaaaggtg 420  
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<400> 30

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 CTCATCTCTA CCCAGCACGA CGAGACAGTC ACCAACGACG AGATTGCTGC TGACCTGAAG  
 GAGCACGTCA TCAAGCCAGT CATCCCCGAG CAGTACCTCG ACGAGAAGAC AATCTTCCAC  
 CTCAACCCGT CTGGCCGNTT CGTCATCGGC GGACCTNACG GCGACGCCGG CCTACTGGCC  
 GNAAGATCAT CATCGACACC TACGGTGGCT GGGGAGCCCA CGGCGGGGGC GCCTTCTTCG  
 GCAAGGACCC GACCAANGTG GACCGCACGG GGCCTACGTN CGAGGNAAGC TTGCNA

ctcacagagg ttgcaagaa tggaacctgg cctgggtca ggcccgatgg gaagaccag 60  
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 ctcatctcta cccagcacga cgagacagtc accaacgacg agattgctgc tgacctgaag 180  
 gagcacgtca tcaagccagt catccccgag cagtacctcg acgagaagac aatcttccac 240  
 ctcaaccctg ctggccgntt cgtcatcggc ggacctnacg gcgacgccgg cctactggcc 300  
 gnaagatcat catcgacacc tacgggtggct ggggagccca cggcgggggc gccttcttcg 360  
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<210> 31  
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<400> 31

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 aacttcgatt tcaggcctgg catgatcatc atcaaccttg acctcaagaa aggcggcaac 360

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<210> 32  
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 ccgtgttcgt cgactcgta cggcaccggca cgatccccga caaggagatc ctcaagtcgt 240  
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 cctgggaagt ggtgaagccc ctcaagttcg acaaggcatc cggcttaang ttggaatnnt 420  
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<210> 33  
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 <213> Zea mays  
 <400> 33

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 tgatggtggt cggcgagatc acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg 360  
 acacctgccg cgagatcggg ttacactccg acgacgtggg cctcgacgcc gaccgctgca 420  
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488

<210> 34  
<211> 480  
<212> nucleic acid  
<213> Zea mays

<400> 34

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gaggtcggta gagcgagaag aaggcaatgg cggccgagag cttccttttc acctcggagt 180  
ccgtgaacga ggggcacccc gacaagctgt gcgaccaggt gtcggacgcc gtgcttgacg 240  
catgcctcgc gcaggacccc gacagcaagg tggcctgcga gacctgcacc aagaccaaca 300  
tggtgatggt gttcggcgag atcacgacca aggcgaccgt ggactacgag aagatcgtgc 360  
gcgacacctg ccgcgagatc gggttcacct ccgacgacgt gggcctcgac gccgaccgct 420  
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<210> 35  
<211> 453  
<212> nucleic acid  
<213> Zea mays

<400> 35

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ccgtgaacga ggggcaccca gacaagctgt gcgaccaggt gtcggacgcg gtgctggacg 180  
cctgcctggc gcangacccc gacagcaagg tggcctgcna gacctgcacc aagacgaaca 240  
tggtgatggt gttcggcgag atcaccacca aggcgagcgt ggactacgag aagatcgtgc 300  
gcgacacctg ccgcgagatc gggttcacct tcgacgacgt ggggctcgac nccnaccgct 360  
gcaaggtgct ggtgaacatc gagcagcagt ccccgacat cgcgcaaggc gtgcacggca 420  
ctttacgaaa ccggcccag gagatcggcc cnt 453

<210> 36  
<211> 505  
<212> nucleic acid



<213> Zea mays

<400> 36

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 gcagcagcgc aagaggttg tagagcgagc gagaagaagg caatggcggc ggagagcttc 180  
 ctgttcacct cggagtccgt gaacgagggg caccagaca agctgtgcga ccaggtgtcg 240  
 gacgcggtgc tggacgcctg cctggcgcan gaccccgaca gcaaggtggc ctgcgagacc 300  
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 tacgagaaga tcgtgcgcga cacctgccgc gagatcgggt tcacctncca cgacgtgggg 420  
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<210> 37

<211> 447

<212> nucleic acid

<213> Zea mays

<400> 37

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 gccctcagc catgtccttg ccaccaagct tgggtgctcgt ctacagagg ttcgcaagaa 180  
 tggaaacctgc ccctggctca ggcccgatgg gaagaccag gtgacagtgg agtaccgcaa 240  
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 cgagacangt caccaacgac gaagattgct gctgacctga aaggaacaac gtcataaac 360  
 caagtcatnc ccgaacagta ctttgacga gaagacaatc tttcanctta accgctctgg 420  
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<210> 38

<211> 420

<212> nucleic acid

<213> Zea mays

<400> 38

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 accaaggttg accgcagcgg agcctatgtc gcaaggcagg ctgccaagag catcgtcgcc 180  
 agcggccttg ctgcgcgcgc catcgtccag gtgtcttacg ccatcggcgt gcccgagcct 240  
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 attgtcaaag gagaacttcg atttcaggcc tggcatgata atcatcaacc ttgacctcaa 360  
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<210> 39  
 <211> 499  
 <212> nucleic acid  
 <213> Zea mays

<400> 39  
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 ggaccaaac aaggttgacc gcagcggagc ctatgtcgca aggcaggctg ccaagagcat 180  
 cgtcgccagc ggcccttgctc gccgcgccat cgtccagggt tcttacgcca tcggcgtgcc 240  
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 cctcaagatt gtcaaggaga acttcgattt caggcctggc atcatcatca accttgacct 360  
 caagaaaggc ggcaacgggc gctacctnaa gacggcggcc tacggcactt tggaaggga 420  
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 gcgagaagaa ggcaatggcg gccgagagct tccttttcac ctcggantcc gtgaacgggg 180



ggacgccggc ctcaccggcc gcaagatcat catcgacacc tacggcggct ggggagccca 300  
cggcgggggc gccttctccg gcaag 325

<210> 43  
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<213> Zea mays  
  
<400> 43

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ccgaggtgcg caagaacggc acctgcgcct ggctgaggcc cgacggcaag acccaggtga 180  
cggtgaggta cgtgaacgag ggcgggcgcca tggtgcccgt ccgcgtgcac accgtgctca 240  
tctccaccca gcacgacgag accgtcacca acgacgagat cgccgccgac ctcaaggagc 300  
acgtcatcaa gcccgtgat 319

<210> 44  
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<212> nucleic acid  
<213> Zea mays  
  
<400> 44

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gagacagtca ccaacgacga gattgctgct gacctgaagg agcacgtcat caagccagtc 180  
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tacggtggct ngggagccca cggcgggggc gccttnttcg gcaaggacc gaccaaagtt 360  
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nccgcggnc 429

<210> 45  
<211> 315  
<212> nucleic acid  
<213> Zea mays

<400> 45

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cctgcgcctg gctgaggccc gacggcaaga ccaggtgac ggtggagtac gtgaacgagg 180

gcggcgccat ggtgcccgtc cgcgtgcaca ccgtgctcat ctccaccag cacgacgaga 240

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ctgagaagta cctcg 315

<210> 46

<211> 474

<212> nucleic acid

<213> Zea mays

<400> 46

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anaannccct agagtnagtc gtttaacggc gggggcgcct tctccggcaa ggacccgacc 120

aaggtggacc gcagcggggc ctacgtcgcg aggcaggctg ccaagagcat cgtcgccgcc 180

ggcctcgccc gccgtgccat cgtccaggtc tctacgcc a tcggcgtgcc cgagcccctg 240

tcggtgttcg tggacacgta cggcaccggc gcgatccccg acaaggagat cctgaagatc 300

gtgaaggaga acttcgactt caggcccggc atgatcatca tcaacctcga cctcaagaaa 360

ggcggcaacg ggcgtacct caagacggcg gcctacgggc actttgggag ggacgaaccc 420

gacttcacct gggaagtngt taaaccccc naaggcgga aanccttntt ctgg 474

<210> 47

<211> 410

<212> nucleic acid

<213> Zea mays

<400> 47

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gggcacttca cgaagcggcc cgaggagatc ggcgcgggcg accagggccca catgttcggg 180

tacnccaccg acnagacccc cgagctgatg ccgctcagcc acgtgctggc caccaagctn 240

ggcncgcgcc tcaccgaggt tccgcaagac gggnacctgc gcctggntga nggccccgacg 300  
gcaagaccaa ggtnacgggtg gagtacgtga actagggcgg ctccattggt gccctccgcg 360  
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<210> 48  
<211> 297  
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<213> Zea mays  
<400> 48

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cacgggggacg ccggcctcac cggccgcaag atcatcatcg acacctacgg cggctgggga 240  
gccacggcg ggggcgcctt ctccggcaag gacccacca aggtggaccg cagcggg 297

<210> 49  
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<212> nucleic acid  
<213> Zea mays  
<400> 49

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ggcaggctgc caagagcatc gtcgccgccc gcctcggccc ccgtgccatc gtccagggtct 180  
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tgatcatcat caacctngac ctcaagaaag gcggnnacgg nccgtacct taaanaacgg 360  
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<211> 316  
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cgacacctgc cgcgagatcg ggttcacctc cgacgacgtg gggctcgcgc ccgaccgctg 240  
caaggtgctg gtgaacatcg ancagcagtc ccccgacntc ggcgagggcg tgcacgggca 300  
nttcacgaag cggccc 316

<210> 51

<211> 339

<212> nucleic acid

<213> Zea mays

<400> 51

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tcttccacct caaccogtct ggccgcttcg tcctcgggcg acctcacggc gacgcggctc 180  
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cttctccggc aangacccga ccaaggtgga ccgcagcggg gcctacgtcg cgaggcaggc 300  
tgccaagagc atcgtcgccg cggcctcgcc gcngcgctt 339

<210> 52

<211> 300

<212> nucleic acid

<213> Zea mays

<400> 52

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gcgtgcacgg gcacttcacg aagcggcccg aggagatcgg cgcgggacgac cagggccaca 180  
tgttcgggta cgccaccgac gagaccccg agctgatgcc gctgagccac gtgctggcca 240  
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<210> 53

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 gacgaccccg acttcacctg ggaggtggtg aagccctca aggcggaaaa acctttctg 420  
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<210> 55  
 <211> 487  
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<210> 58

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 gtacgtgaac ganggcggcg ccatggtgcc cgtccgctg cacaccgtgc tcctctccac 180  
 ccagcacgac gagaccgtca ccaacgacga gatcgccgcc gacctcaagg agcacgtcat 240  
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<210> 60  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 60  
  
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 ctaggtgctc gtctcaccga ggtccgcaag aacggaacct gccctggct caggcctgat 120  
 gggaagaccc aggtgacagt cgagtaccgc aatgaggggtg gtgccatggt ccccatccgt 180  
 gtccacaccg tctcatctc caccagcac gacgagacag tgaccaatga tgagatcgct 240  
 gctgacctga aggagcatgt catcaagcct gtcacccctg agcagtacct tgacgagaag 300

accat

305

<210> 61  
<211> 300  
<212> nucleic acid  
<213> Zea mays  
  
<400> 61

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ggcgaccgtg gactacgaga agatcgtgcg cgacacctgc cgcgagatcg ggttcacctc 120  
cgncgacgtg ggcctcgacg ccgnccgctg caaggtgctg gtgaacatcg agcagcagtc 180  
ccccgacatc gcgcagggcg tgcacgggca cttcacgnag cggnccgagg agatnggngc 240  
gggcgaccag ggncacatgt tcgggtacgn caccgacgag acccccgagc tgatgccgct 300

gagcagcagc gcaagaggtc ggtagagcga gaagaaggca atggcgggccg agagcttcct  
tttcacctcg gaggccgtga acgaggggca ccccgacaag ctgtgcgacc aggtgtcgga  
cgccgtgctt gacgcatgcc tcgcgcagga ccccgacagc aaggtggcct gcgagacctg  
caccaagacc aacatggtga tgggtgttcg cgagatcacg accaaggcga ccgtggacta  
cgagaagatc gtgcgcgaca cctgccgcga gatcgggttc acctncgacg acgtgggcct  
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anggcgttca cnggcant

<210> 62  
<211> 558  
<212> nucleic acid  
<213> Zea mays  
  
<400> 62

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ggaattcccg ggtcgaccca cgcgtccgcc cagcgtccg cccacgcgtc cgcccacgog 120  
tccgcccacg cgtcgcgcca cgcgtccggc ctcttctccc tcttgccggt cccgaataaa 180  
gagcagcagc gcaagaggtc ggtagagcga gaagaaggca atggcgggccg agagcttcct 240  
tttcacctcg gaggccgtga acgaggggca ccccgacaag ctgtgcgacc aggtgtcgga 300  
cgccgtgctt gacgcatgcc tcgcgcagga ccccgacagc aaggtggcct gcgagacctg 360  
caccaagacc aacatggtga tgggtgttcg cgagatcacg accaaggcga ccgtggacta 420  
cgagaagatc gtgcgcgaca cctgccgcga gatcgggttc acctncgacg acgtgggcct 480  
tnacgcnac cnntgcaagg tncgtgtgaa cattgagcaa naattcccng gactttnnge 540  
anggcgttca cnggcant 558

<210> 63  
<211> 332  
<212> nucleic acid  
<213> Zea mays

<400> 63

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 gtgaccaatg atgagatcgc tgctgacctg aaggagcatg tcatcaagcc tgtcatccct 180  
 gagcagtacc ttgacgagaa gaccatcttc caccttaacc catctggccg ctttgtcatg 240  
 tggacctcac ggcgatgctg gcctcactgg ccgcaagatc atcatgacac ctacgggtggc 300  
 tggggagccc atggtggtgg cgctttctcc gg 332

<210> 64

<211> 314

<212> nucleic acid

<213> Zea mays

<400> 64

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 gaccaacatg gtcatggtct ttggtgagat caccaccaag gccaatgttg actacgagaa 180  
 gattgtcagg gagacctgcc gcaacattgg ttttgtgtca aacgatgttg ggcttgacgc 240  
 cgaccactgc aagtgtctgt gaacattnag cagcagtcct ctgatattgc tcanggtgtg 300  
 catggccact tcac 314

<210> 65

<211> 293

<212> nucleic acid

<213> Zea mays

<400> 65

ggacgagaag accatcttcc acctcaaccc gtcggggcgc ttctcatcgc gcgggcccc 60  
 cggggacgcc ggctcaccg gccgcaagat catcatcgac acctacggcg gctggggagc 120  
 ccacggcggg ggcccttctt ccggcaagga cccaccaag gtggaccgca gcggggccta 180  
 cgtcgccagg caggccgcca agagcatcgt ggccagcggc ttgcccgcgc gctgcctcgt 240  
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<210> 66

<211> 289  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 66  
  
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 ccaccaaggc gagcgtggan tacgagaagn tcgtgcgcga cacctgccgc gagatcgggt 120  
 tcacctccga cgacgtgggg ctgcacgccg accgctgcaa ggtgctggtg aacatcgagc 180  
 agcagtcccc cgacatcgcg cagggcgtgc acgggcactt cacgaagcgg cccgaggaga 240  
 tcggcgccgg cgaccagggc cacatgttcg ggtacgccac cgacgagac 289

<210> 67  
 <211> 306  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 67  
  
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 gttgggcttg acgccgacca ctgcaagggtg ctctgaaca ttgagcagca gtcccctgat 120  
 attgctcagg gtgtgcatgg ccacttcacc aagcgccccg aggagattgg agctggtgac 180  
 caggacaca tgttcgggta tgcgaccgat gagaccctg agttgatgcc cctcagccat 240  
 gtcttgcca ccaagctagg tgctcgtctc accgaggtcc gcaagaacgg aacctgcccc 300  
 tggctc 306

<210> 68  
 <211> 303  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 68  
  
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 tgacagcaag gttgcttggtg agacctgcac caagaccaac atggtcatgg tctttggtga 120  
 gatcaccacc aaggccaatg ttgactacga gaagattgtc agggagacct gccgcaacat 180  
 tggttttgtg tcaaacgatg ttgggcttga cgccgaccac tgcaagggtgc tcgtgaacat 240  
 tgagcagcag tcccctgata ttgctcaggg tngcatggc cacttcacca agcgccccga 300

gga

303

<210> 69  
<211> 300  
<212> nucleic acid  
<213> Zea mays

<400> 69

caaagaccaa catggtcatg gtcttttggtg agatcaccac caaggccaat gtcgactacg 60  
agaagattgt cagggagaca tgccgcaaca ttggtttcgt ntcgaacgat gtcgggcttg 120  
acgtgacca ctgcaagggtg cttgtgaaca ttgagcagca gtcccctgat attgctcagg 180  
gtgtncacgg ccacttcacc aagcgccccg aggagattgg agctggtgac caggggcaca 240  
tgtttgggta tgcgactgac gagaccctg agctgatgcc cctcagccat gtccttgcca 300

<210> 70  
<211> 329  
<212> nucleic acid  
<213> Zea mays

<400> 70

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gacaccctga caagctctgt gaccaggtct cagatgccgt tcttgacgct tgccttgctg 120  
aggaccctga cagcaagggtt gcttgtgaga cctgcaccaa gaccaacatg gtcattggtct 180  
ttggtgagat caccaccaag gccaatgttg actacgagaa gattgtcagg gagacctgcc 240  
gcaacattgg ttttgtgtca aacgatgttg ggcttgacgc cgaccattgc aaggtgcncg 300  
tgaanatnng cancagtcct ctgatattg 329

<210> 71  
<211> 304  
<212> nucleic acid  
<213> Zea mays

<400> 71

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gtcttttgng agatcaccac caaggccaat gtcgactacg agaagattgt cagggagaca 120  
tgccgcaaca ttggtttcgt gtcgaacgan gtcgggcntg angctgacca ctgcaaggng 180

cttgtgaaca ttgagcagca gtccccctgat attgctcagg gtgtgcacgg ccacttcacc 240  
aagcgccccg aggagattgg agctggtgac caggggcaca tgtttgggta tgcgactnac 300  
gaga 304

<210> 72  
<211> 307  
<212> nucleic acid  
<213> Zea mays

<400> 72

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catggtcang gnccttttggg gagatcacca ccaaggccaa tgttgactac gagaagattg 120  
tcagggagac ctgccgcaac attggttttg tgtcaaacga tgttgggctt gacgccgacc 180  
actgcaaggt gctcgtgaac attgagcagc agtccccctga tattgctcag ggtgtgcatg 240  
gccacttcac caagcgcccc gaggagattg gagctggtga ccagggacac atgttcgggt 300  
atgcgac 307

<210> 73  
<211> 282  
<212> nucleic acid  
<213> Zea mays

<400> 73

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gcgcgcctga cggaggtccg caaggacggc acctgcgcct ggctcaggcc cgacggcaag 120  
accaggtga cgggtggagta cgtgaacgag ggcggcgcca tgggtcccgt ccgcgtgcac 180  
accgtgctca tctccaccca gcacgacgag accgtcacca acgacgagat cgccgccgat 240  
ctcaaggagc acgtcatcaa gcccgtcac ccggagaggt ac 282

<210> 74  
<211> 320  
<212> nucleic acid  
<213> Zea mays

<400> 74

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cgatgggaag acccaggtga cagtggagta ccgcaacgag ggtggcgcca tggttcccat 120  
ccgtgtgcac acagtcctca tctctacca gcacgacgag acagtcacca acgacgagat 180  
tgctgctgac ctgaaggagc acgtcatcaa gccagtcac cccgagcagt acctcgacga 240  
gaagacaatc ttccacctca acccgtctgg ccgcttcgtc atcggcggac tcacggcgac 300  
ctggcctcac tggccggaag 320

<210> 75  
<211> 370  
<212> nucleic acid  
<213> Zea mays  
<400> 75

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gcgtgccgga gccctgtca gtgttcgtcg actcctacgg caccgggacc atccccgaca 120  
aggagatcct caagatcgtc aaggagaact tcgacttcag gcccgggatg atcaccatca 180  
acctcgacct caagaagggc ggcaacaggt tcatcaagac cgccgcatac ggccaacttg 240  
gccgtgacga cgccgacttc acctgggagg tggtaagcc cctaaagaag gcatccgctt 300  
aagaatgtat tgggaagttc actggacatg aggttcactc tcgtctggct ctgctgatac 360  
ctgcaaggat 370

<210> 76  
<211> 300  
<212> nucleic acid  
<213> Zea mays  
<400> 76

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acgaggggtg cgccatgggt cccatccgtg tgcacacagt cctcatctct acccagcacg 120  
acgagacagt caccaacgac gagattgctg ctgacctgaa ggagcacgtc atcaagccag 180  
tcatccccga gcagtacctc gacgagaaga caatcttcca cctcaacccg tctggccgct 240  
tcgtcatcgg cggacctcac ggcgacgctg gcctcactgg ccggaagatc atcatcgaca 300

<210> 77  
<211> 315  
<212> nucleic acid



<213> Zea mays

<400> 77

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gacctgcacc aagaccaaca tggatcatggt ctttggtgag atcaccacca aggccaatgt 120  
tgactacgag aagattgtca gggagacctg ccgcaacatt ggttttgtgt caaacgatgt 180  
tgggcttgac gccgaccact gcaaggtgct cgtgaacatt gagcagcagt cccctgatat 240  
tgctcagggt gtgcatggcc attcaccaag cgccccgang agattggagc tggtgaccag 300  
gacacatggt cgggg 315

<210> 78

<211> 297

<212> nucleic acid

<213> Zea mays

<400> 78

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tgcaccaaga ccaacatggt catggtcttt ggtgagatca ccaccaaggc caatgttgac 180  
tacgagaaga ttgtcagggg gacctgccgc aacattgggt ttgtgtcaaa cgatgttggg 240  
cttgacgcgc accactgcaa gtgctcgtga acattgagca gcagtcccct gatattg 297

<210> 79

<211> 448

<212> nucleic acid

<213> Zea mays

<400> 79

ttggttnagat caccaccaag gccaatgttg actacganaa gattgtnagg ganacctgtc 60  
gnacattgg ttttgtgtca aacgatgttg ggcttgacgc cgaccactgc aagggtgctcg 120  
tgaacattna gcagnagtnc cctgatattg ctanggtgt gcatggccac ttnaccaanc 180  
gccccganga gattgganct ggtgaccagg gacacatggt cgggtatgcg accgatgaga 240  
cccctnagtt gatgccctc agccatgtcc ttgccaccaa gctaggtgct cgtctnaccg 300  
aggtncncaa gaaccggaac ctgccnctgg ctangcctg atgngaagac cnatgtgaca 360

gtcnantnnc gnaatgaagg gtggtgccat tgnccccatc ctngtcaaca ccgttcttat 420  
 ttcaaccaag tnngacgagg acaatgac 448

<210> 80  
 <211> 287  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 80

caccgtcctc atctccaccc agcacgacga gacagtgacc aatgatgaga tcgctgctga 60  
 cctgaaggag catgtcatca agcctgtcat cctgagcag taccttgacg agaagaccat 120  
 cttccacctt aacccatctg gccgctttgt cattgggtgga cctcacggcg atgctggcct 180  
 cactggccgc aagatcatca ttgacaccta cgggtggctgg ggagcccatg gtggtggcgc 240  
 tttctccggc aaggacccaa ccaaggttga ccgcagcgga gctatgt 287

<210> 81  
 <211> 290  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 81

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 acgacgagac agtgaccaat gatgagatcg ctgctgacct gaaggagcat gtcacaaagc 120  
 ctgtcatccc tgagcagtac cttgacgaga agaccatctt ccaccttaac ccatctggcc 180  
 gctttgtcat tgggtggacct cacggcgatg ctggcctcac tggccgcaag atcatcattg 240  
 acacctacgg tggctgggga gcccatggtg gtggcgcttt ctccggcaag 290

<210> 82  
 <211> 287  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 82

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 cgaacatggt gatggtgttc ggcgagatca ccaccaaggc gagcgtggac tacgagaaga 120  
 tcgtgcgcga cacctgccgc gagatcgggn nnacctccga cgacgtgggg ctgcacgccg 180

accgctgcaa ggtgctggtg aacatcgagc agcagtcgcc cgacatcgcg cagggcgtgc 240  
acgggcactt cacgaagcgg cccgaggaga tcggcgccgg cgaccag 287

<210> 83  
<211> 291  
<212> nucleic acid  
<213> Zea mays  
  
<400> 83

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aacggaacct gccctggct caggcctgat gggaagacct aggtgacagt cgagtaccgc 120  
aatgaggggtg gtgccatggt ccccatcgt gtccacaccg tcctcatctc caccagcac 180  
gacgagacag tgaccaatga tgagatcgct gctgacctga aggagcatgt catcaagcct 240  
gtcatccctg agcagtacct tgacgagaag accatcttcc accttaacct a 291

<210> 84  
<211> 283  
<212> nucleic acid  
<213> Zea mays  
  
<400> 84

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acggcacctg cgcttggtg aggcccgatc ggcaagacct aggtgacggt ggagtacgtg 120  
aacgagggcg ggcctatggt gccgctccgc gtgcacaccg tgctcatctc caccagcac 180  
gacgagaccg tcaccaacga cgagatcgcc gccgacctca aggagcacgt catcaagccc 240  
gtgatccctg agaagtacct cgacgagaag accatcttcc acc 283

<210> 85  
<211> 274  
<212> nucleic acid  
<213> Zea mays  
  
<400> 85

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atgcctcgcg caggaccccg acagcaaggt ggctgcgag acctgcacca agaccaacat 120  
ggtgatggtg ttcggcgaga tcacgaccaa ggcgacctg gactacgaga agatcgtgcg 180

cgacacctgc cgcgagatcg ggttcacctc cgacgacgtg ggctcgcaca ccgaccgctg 240  
caaggtgctg gtgaacatcg agcagcagtc cccc 274

<210> 86  
<211> 290  
<212> nucleic acid  
<213> Zea mays  
<400> 86

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cctcaaccog tctggccgct tcgtcatnng cggacctcac ggcgacgctg gcctcaactgg 120  
ccggaagatc atcatcgaca cctacgggtgg ctgggggagcc cacggcgggg gcgccttctc 180  
cggcaaggac ccgaccaagg tggaccgcag cgggggctac gtcgcgaggc aggttgccaa 240  
gagcatcgtc gccgcgggcc tcgccgcng tgccatcgtc caggtctcct 290

<210> 87  
<211> 290  
<212> nucleic acid  
<213> Zea mays  
<400> 87

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gaccaggtct cagatgccgt tcttgacgct tgccttgctg aggaccctga cagcaagggt 120  
gcttgtgaga cctgcaccaa gaccaacatg gtcattgtct ttggtgagat caccaccaag 180  
gccaatgttg actacgagaa gattgtcagg gagacctgcc gcaacattgg ttttgtgtca 240  
aacgatgttg ggcttgacgc cgaccactgc aaggtgctcg tgaacattga 290

<210> 88  
<211> 288  
<212> nucleic acid  
<213> Zea mays  
<400> 88

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tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc cgacagcaag 120  
gtggcctgcg agacctgcac caagaccaac atggtgatgg tggtcggcga gatcacgacc 180

aaggcgaccg tggactacga gaagatcgtg cgcgacacct gccgcgagat cgggttcact 240  
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<210> 89  
 <211> 289  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 89

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 gaaggagcac gtcataaagc cagtcacccc cgagcagtag ctcgacgaga agacaatctt 180  
 ccacctcaac ccgtctggcc gcttcgtcat cggcggacct cacggcgacg ctggcctcac 240  
 tggccggaag atcatcatcg acacctacgg tggctgggga gccacggc 289

<210> 90  
 <211> 330  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 90

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 gcaatggcgg ccgagagctt ccttttcacc tcggagtccg tgaacgaggg gcaccccgac 120  
 aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180  
 agcaagggtgg cctgcgagac ctgcaccaag accaacaatgg tgatggtgtt cggcgagatc 240  
 acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acacctgccg cgagatcggg 300  
 ttcacctccg acgacgtggg cctcgacgcc 330

<210> 91  
 <211> 291  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 91

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 agaccaggt gacagtcgag taccgcaatg aggggtgtgc catggtcccc atccgtgtcc 120

acaccgtcct catctccacc cagcacgacg agacagtgac caatgatgag atcgctgctg 180  
acctgaagga gcatgtcatc aagcctgtca tccctgagca gtaccttgac gagaagacca 240  
tcttccacct taacccatct ggccgctttg tcattggtgg acctcacggc g 291

<210> 92  
<211> 285  
<212> nucleic acid  
<213> Zea mays

<400> 92

gccagcggcc tgcgccgcg ctgcctcgtg caggtgtcct acgccatcgg cgtgccggag 60  
cccctgtccg tgttcgtcga ctctacggc accgggacca tccccgacaa ggagatccta 120  
aagatcgtca aggagaactt cgacttcagg ccagggatga tcaccatcaa cctcgacctc 180  
aagaagggcg gcaacagggt catcaagacc gccgcatacg gccactttgg ccgtgacgac 240  
gccgacttca cctgggagggt ggtcaagccc ctaaagaagg catcc 285

<210> 93  
<211> 283  
<212> nucleic acid  
<213> Zea mays

<400> 93

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tatgcgaccg atgagacccc tgagttgatg ccctcagcc atgtccttgc caccaagcta 120  
ggtgctcgtc tcaccgaggt ccgcaagaac ggaacctgcc cctggctcag gcctgatggg 180  
aagacccagg tgacagtcga gtaccgcaat gaggggtgggt ccatggtccc catccgtgtc 240  
cacaccgtcc tcatctccac ccagcacgac gagacagtga cca 283

<210> 94  
<211> 298  
<212> nucleic acid  
<213> Zea mays

<400> 94

actacgagaa gattgtcagg gagacatgcc gcaacattgg ttctgtgtcg aacgatgtcg 60  
ggcttgacgc tgaccactgc aaggtgcttg tgaacattga gcagcagtc cctgatattg 120

ctcaggggtgt gcacggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 180  
ggcacatggt tgggtatgcn actgacgaga cccttgagct gatgcccctc agccatgtcc 240  
ttgccaccaa gcttgggtgtc gtctcacnga aggttcgcaa gaatggaacc tgcccctt 298

<210> 95  
<211> 469  
<212> nucleic acid  
<213> Zea mays

<400> 95

cttctccctc ttgccgggtcc cgaataaaga gcagcagcgc aagagggtcgg tagagcgaga 60  
agaaggcaat ggcggccgag agcttctttt tcacctcgga gtccgtgaac gaggggcacc 120  
ccgacaagct gtgcnaccag gtgtcggacg ccgtgcttga cncatgcctc gcgcaggacc 180  
ccnacaccaa ggtggcctgc nagacctgca ccaanaccaa catggtgatg gtgttcggcg 240  
agatcacgac caangcgacc gtggactacg agaagatcgt gcgccgacac ctgccgag 300  
atcggttca ccttcgncga cgtgngccct tgactccnnc ccggtgcaag gtgctggtga 360  
acattnatca tcaatncccc gacattnttc aaggcttca cggcacttta cgaaacggcc 420  
cnangagatc ggccggggcca acagngccac atnttcgggt ccccccca 469

<210> 96  
<211> 293  
<212> nucleic acid  
<213> Zea mays

<400> 96

aacgatgttg ggcttgacgc cgaccactgc aaggtgctcg tgaacattga gcagcagtcc 60  
cctgattgct caggggtgtgc atggccactt caccaagcgc cccgaggaga ttggagctgg 120  
tgaccaggga cacatgttcg ggtatgcnac cgatgagacc cctgagttga tgcccctcag 180  
ccatgtcctt gccaccaagc taggtgctcg tctcaccgag gtccgcaaga acggaacctg 240  
cccctggctc aggctgatg ggaagacca ggtgacagtc gagtaccgca aaa 293

<210> 97  
<211> 280  
<212> nucleic acid  
<213> Zea mays

<400> 97

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gtgctggacg cctgcctggc gcagganccc gacagcaagg tggcctgcga gacctgcacc 120

aagacgaaca tggatgatggg gttcggcgag atcaccacca aggcgagcgt ggactacgag 180

aagatcgtgc gcgacacctg ccgcgagatc gggttcacct ccgacgacgt ggggctcgac 240

gccgaccgct gcaagggtgct ggtgaacatc gagcagcagt 280

<210> 98

<211> 285

<212> nucleic acid

<213> Zea mays

<400> 98

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gtcgcgaggc aggttgccaa gagcatcgtc gccagcggcc ttgctcgccg cgccatcgtc 120

cagggtgtctt acgccatcgg cgtgcccagag cctctctccg tgttcgtcga cacgtacggc 180

accggcgcga tccccgacaa ggagatcctc aagattgtca aggagaactt cgatttcagg 240

cctggcatga tcatcatcaa ccttgacctc aagaaaggcg gcaag 285

<210> 99

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 99

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tcctcatctc caccagcac gacgagacag tgaccaatga tgagatcgct gctgacctga 120

aggagcatgt catcaagcct gtcatncctg agcagtacct tgacgagaag accatcttcc 180

accttaacct atctggccgc tttgtcattg gtggacctca cggcgatgct ggctcactg 240

gccgcaagat catcattgac acctacggtg gctgggga 278

<210> 100

<211> 275

<212> nucleic acid

<213> Zea mays



<400> 100

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gagcagtacc ttgacgagaa gaccatcttc caccttaacc catctggccg ctttgtcatt 120

gggtggacctc acggcgatgc tggcctcact ggccgcaaga tcatcattga cacctacggt 180

ggctgggggag cccatgggtgg tggcgctttc tccggcaagg acccaaccaa ggttgaccgc 240

agcggancct atgtcgcaag gcaggctgcc aagag 275

<210> 101

<211> 291

<212> nucleic acid

<213> Zea mays

<400> 101

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cgagaagacc atcttccacc ttaaccctac tggccgcttt gtcattgggtg gacctcacgg 120

cgatgctggc ctactggcc gcaagatcat cattgacacc tacggtgggt ggggagccca 180

tggtgggtggc gctttctccg gcaaggaccc aaccaagggt gaccgcagcg gaggcctatgt 240

cgcaaggcag gctgccaaga gcatcgtcgc cagcggcttg ctgcccgcgc c 291

<210> 102

<211> 301

<212> nucleic acid

<213> Zea mays

<400> 102

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ctgacaagct ctgagaccag gtctcagatg ctgttctgga cgcttgccctg ctgaggaccc 120

tgacagcaag gttgcttgcg agacctgcac caagaccaac atgggtcatgg tctttggtga 180

gatcaccacc aaggccaatg tcgactacga gaagattgtc agggagacat gccgcaacat 240

tggtttcgtg tcgaacgatg tcgggcttga cgctgaccac tgcaagtgt tgtgaacatt 300

g 301

<210> 103

<211> 336

<212> nucleic acid

<213> Zea mays

<400> 103

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aggcaatggc ggccgagagc ttcttttttna cctcggagtc cgtgaaacga ggggcncccc 120  
gacaagctgt gcgaccaggt gtcggacgcc gtgcttgacg catgcctcgc gcaggacccc 180  
gacagcaagg tggcctgcga gacctgcacc aagaccaaca tggatgatggt gttcggcgag 240  
atcacgacca aggcgaccgt ggactacgag aagatcgtgc gcgacacctg ccgcgagatc 300  
gggttcacct ccgacgacgt gggcctcgac gccgac 336

<210> 104

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 104

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cggcggggccc cagcgtgacg ccggcctcac cggccgcaag atcatcatcg acacgtacgg 120  
cggctgggga gcccacggcg gtggcgccctt ctccggcaag gacccacca aggtggaccg 180  
cagcggcgcc tacgtggcca ggcaggccgc caagagcatc gtggccagcg gctcgcccgc 240  
cgctgcctcg tgcagtgtcg tacgccatcg ctgccg 276

<210> 105

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 105

gagattggag ctggtgacca gggacacatg ttcgggtatg cgaccgatga gacccttgag 60  
ttgatgcccc tcagccatgt ccttgccacc aagctaggtg ctggtctcac cgaggctcgc 120  
aagaacggaa cctgcccctg gctcaggcct gatgggaaga cccagggtgac agtcgagtac 180  
cgcaatgagg gtggtgccat ggtccccatc cgtgtccaca ccgtcctcat ctccaccag 240  
cacgacgaga cagtgaccaa tgatgagatc gtgctgacct gaaggag 287

<210> 106

<211> 303  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 106  
  
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 ctgaaggagc atgtcatcaa gcctgtcatc cctgagcagt accttgacga gaagaccatc 120  
 ttccacctta acccatctgg ccgctttgtc attggtggac ctcacggcga tgctggcctc 180  
 actggccgca agatcatcat tgacacctac ggtggctggg gagcccatgg tgggtggcgct 240  
 ttctccggca aggacccaac caagttgacc gcagcgganc tatgtcgcaa ggcagctgcc 300  
 aag 303

<210> 107  
 <211> 279  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 107  
  
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 ttgcttgtga gacctgcacc aagaccaaca tggatcatgt ctttggtgag atcaccacca 120  
 aggccaatgt tgactacgag aagattgtna gggagacctg ccgcaacatt ggttttgtgt 180  
 caaacgatgt tgggcttgac gccgaccact gcaagggtgt cgtgaacatt gagcagcagt 240  
 cccctgatat tgctcagggt gtgcatggcc acttcacca 279

<210> 108  
 <211> 330  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 108  
  
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 gaagaaggca atggcggccg agagcttcct tttcacctcg gagtccgtga acgaggggca 120  
 ccccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc tcgcgcagga 180  
 ccccgacagc aagggtggcct gcgagacctg caccaagacc aacatgggtga tgggtgttcgg 240  
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gatcgggttc acctccgacg acgtgggcct

330

<210> 109  
<211> 298  
<212> nucleic acid  
<213> Zea mays

<400> 109

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tccgntgca caccgtgctc atctccaccc agcacgacga gaccgtcacc aangacgagn 120  
tcgcccgcgn cctcanggag cacgtcntna agcccgatcat cccgganagg tacctggacg 180  
anaagacnt cttnacctc aaccggtcgg gggcgnttcg tctcggcgcg gcccacggg 240  
gacnccggcc tnaccggccg caagntgntc ntngncacct acngcggntg gggagccc 298

<210> 110  
<211> 498  
<212> nucleic acid  
<213> Zea mays

<400> 110

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cccacggatt ttggntctn ctccggcaag gacccaccn nggtgggggn gnattgggnc 120  
ctaccgtcgc caggcangcc gacaagagca tngnggccag cggcctcgn cgcgntgcc 180  
tcngncaggt gtctacgcc atcggcgtgc cggagcccct gtccgtgttc gtngactcct 240  
acggcaccgg gaccatcccc gacaaggaga tcctaaagat cgtnaaggag aacttcgact 300  
tcaggccagg gatggtcacc atcaacctcg acctcaagaa gggcggaac aggttcatca 360  
agaccgcgn atacggccac tttggcccgt gacgacgccg acttcacctg ggaggtggtc 420  
aagcccctaa agaaggcatc cgcttaagaa tgtattnga aagttcactg gacatgaagg 480  
atcatcttcc tctnggct 498

<210> 111  
<211> 284  
<212> nucleic acid  
<213> Zea mays

<400> 111

gccgcagat caaagaagat ggcagctgtc gacacattcc tttcacctc ggagtctgtg 60  
 aacgagggac accctgacaa gctctgtgac caggtctcag atgccgttct tgacgcttgc 120  
 cttgctgagg accctgacag caaggttgct tgtgagacct gcaccaagac caacatggtc 180  
 atggtctttg gtgagatcac caccaaggcc aatgttgact acgagaagat tgtcagggag 240  
 acctgccgca acattggttt tgtgtcaaac gatgttgggc ttga 284

<210> 112  
 <211> 328  
 <212> nucleic acid  
 <213> Zea mays

<400> 112

ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga cacctgccgc 60  
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 aacatcgagc agcagtcctc cgacatcgcg cagggcgctgc acgggcactt cacgaagcgg 180  
 cccgaggaga tcggcgcggg nnaccagggc cacatgttcg ggtacgccac cgacgagacc 240  
 cccgagctga tgccgctgag ccaacgtgct ggccaacaag ctgggcgcgcg ggctcaccga 300  
 ngtgcgaaaa acggcaactg cgctggct 328

<210> 113  
 <211> 287  
 <212> nucleic acid  
 <213> Zea mays

<400> 113

gggggcgctt tctccggcaa ggacccgacc aaggtggacc gcagcggggc ctacgtcgcg 60  
 aggcaggctg ccaagagcat cgtcgccgcc ggctcgcgcc gccgcgccat tgtccaggtc 120  
 tctacgcca tcggcgtgcc cgagccctt tcggtgttcg tggacacgta cggcaccggc 180  
 gccatccccg acaaggagat cctgaagatc gtgaaggaga acttcgactt caggccccgc 240  
 atgatcatca tcaacctcga cctcaagaaa ggcggcaacg ggcgcta 287

<210> 114  
 <211> 261  
 <212> nucleic acid  
 <213> Zea mays

<400> 114

cgacgccgaa ccgctgcaag gtgctggtga acatcgagca gcagtcccc gacatcgcg 60

agggcggtgca cgggcacttc acgaagcggc ccgaggagat cggcgcgggc gaccagggcc 120

acatgttcgg gtacgccacc gacgagaccc ccgagctgat gccgctgagc cacgtgctgg 180

ccaccaagct gggcgcgcg ctcaccgagg tgcgcaagaa cggcacctgc gcctggctga 240

ggcccgacgg caagaccag g 261

<210> 115  
 <211> 294  
 <212> nucleic acid  
 <213> Zea mays

<400> 115

gggccacttc accaagcgcc ccgaggagat tggagctggt gaccagggac acatgttcgg 60

gtatgcgacc gatgagaccc ctgagttgat gccctcagc catgtccttg ccaccaagct 120

aggtgctcgt ctcaccgagg tccgcaagaa cggaanctgc ccctggctca ggctgatgg 180

gaagaccag gtgacagtcg agtaccgcaa tgagggtggt gccatggtcc ccatccgtgt 240

ccacaccgtc ctcatctcca cccagcacga cgagacatga ccaatgatga gatc 294

<210> 116  
 <211> 318  
 <212> nucleic acid  
 <213> Zea mays

<400> 116

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ttccacctca acccgctctgg ncgcttcgtc atcggcggac ctcacggcga cgccggcctc 120

actggccgga agatcatcat cgacacctac ggtggctggg gagcccacgg cgggggccc 180

ttctccggca aggacccgac caangtgga cgcagcgggg cctacgtcgc gangcaggct 240

gccaagagca tcgtcgccgc cggcctcgcc gcngcgccat cgtccaggtc tctagcatgg 300

gtgccgancc tatcgtgt 318

<210> 117  
 <211> 256  
 <212> nucleic acid

<213> Zea mays

<400> 117

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 ggctggggag cccacggcgg tggcgcttc tccggcaagg accccaccaa ggtggaccgc 180  
 agcggcgctt acgtggccag gcaggccgcc aagagcatcg tggccagcgg cttcgccgcg 240  
 cgctgccttc tgcaag 256

<210> 118

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 118

gtcgacacat tcctcttcac ctgggagtct gtgaacgagg gacaccctga caagctctgt 60  
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 gcttgtgaga cctgcaccaa gaccaacatg gtcatggtct ttggtgagat caccaccaag 180  
 gncnatgttg actacgagaa gattgtcagg gagacctgcc gcaacattgg ttttgtgtca 240  
 aacgatgttg ggcttgacgc cgaccactgc aaggt 275

<210> 119

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 119

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 gcttgtgaga cctgcaccaa gaccaacatg gtcatggtct ttggtgagat caccaccaag 180  
 gccattgttg actacgagaa gattgtcagg gagacctgcc gcaacattgg ttttgtgtca 240  
 aacgatgttg ggcntgacgc cgaccactgc aaggtg 276

<210> 120

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 120

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cctgagcagt accttgacga gaagaccatn ttccacctta acccatctgg ccgctttgtc 120  
attggtggac ctcacggcga tgctggcctc actggccgca agatcatcat tgacacctac 180  
ggtggctggg gagcccatgg tgggtggcgt ttctccggca aggacccaac caaggttgac 240  
cgcagcggag cctatgtcgc aangcangct gccaaagagca tcgtcgccaa cggcttgctc 300  
gccgcgcca 309

<210> 121

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 121

ctcagggtgt gcatggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 60  
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ttgccaccaa gctaggtgct cgtctcaccg aggtccgcaa gaacggaacc tgcccctggc 180  
tcaggcctga tgggaagacc caggtgacag tcgagtaccg caatgagggt ggtgccatgg 240  
tccccatccg tgtccacacc gtctctca 267

<210> 122

<211> 277

<212> nucleic acid

<213> Zea mays

<400> 122

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acttcaccaa gcgccccgag gagattggag ctggtgacca ggggcacatg tttgggtatg 120  
cgactgacga gacccttgag ctgatgcccc tcagccatgt ccttgccacc aagcttggtg 180  
ctcgtctcac ggaggttcgc aagaatgaa cctgcccctg gtcaggccc gatgggaaga 240  
cccagtgaca attggantac cgcaacgagg gtggccc 277

<210> 123



<211> 264  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 123  
  
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 tgaggccccga cggcaagacc caggtgacgg tggagtacgt gaacgagggc ggcgccatgg 180  
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 acgagatcgc ccgccgacct caag 264

<210> 124  
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<210> 125  
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 nacgagggcg gcgccatggt gcccgtccgc gtgcacaccg tgctcatctc cacncancan 180  
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<210> 126

<211> 260  
 <212> nucleic acid  
 <213> Zea mays  
  
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 caacattggt ttogtgtcga acgatgtcgg gcttgacgt gaccactgca aggtgcttgt 180  
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<210> 127  
 <211> 516  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 127  
  
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 cacgtacggc ggctggggag cccacggcgg tggcgcttc tccggcaagg accctaccaa 240  
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 gcctcncctg ccgctgcttc gtgcaggtgt cgtacgcnat cggcgtgcac ggagcccntg 360  
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<210> 128  
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ccgcaagatc atcattgaca cctacggtgg ctggggagcc catggtggtg gcgctttctc 180  
 cggcaaggac ccaaccaagg ttgaccgcag cggagcctat gtcgcaangc aggctgccaa 240  
 gagcatcgtc gccagcggcc ttgc 264

<210> 129  
 <211> 270  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 129

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 gcacgacgag acagtcacca acgacgagat tgctgctgac ctgaaggagc acgtcatcaa 180  
 gccagtcate cccgagcagt acctcgacga gaagacaatc ttccacctca acccgtctgg 240  
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<210> 130  
 <211> 249  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 130

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 aagatcatca tcgacacgta cggcggctgg ggagcccacg gcggtggcgc cttctccggc 180  
 aaggacccca ccaaggtgga ccgcagcggc gcctacgtgg ccaggcaggc cgccaagagc 240  
 atcgtggcc 249

<210> 131  
 <211> 270  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 131

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 ccattctggcc gctttgttat tgggtggacct cacggcgatg ctggcctcac tggccgcaag 120

atcatcattg acacctacgg tggctgggga gcccatgggtg gtggcgcttt ctccggcaag 180  
gacccaacca aggttgaccg cagcggagcc tatgtcgcaa ggcaggctgc caagagcatc 240  
gtcgccagcg gccttgctcg ccgcgccatc 270

<210> 132  
<211> 265  
<212> nucleic acid  
<213> Zea mays

<400> 132

ctgcancaag accaacaatgg tcatggtctt tggtagatc accaccaagg ccaatgttga 60  
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gcttgacgcc gaccactgca aggtgctcgt gaacattgag cagcagtccc ctgatattgc 180  
tcaggggtgtg canggccact tcaccaagcg ccccgaggag attggagctg gtgaccaggg 240  
acacatgttc gggatatgca ccgat 265

<210> 133  
<211> 284  
<212> nucleic acid  
<213> Zea mays

<400> 133

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ccaagagcat cgtngccagc ngcctcgccc gccgctgcct cgtgcagggtg tcgtacgcca 180  
toggntgccg gagncctgt ccgtgttcgt caactcgtac ggcaccggca cgatccccga 240  
caaggagatc ctcaagatcg tgaaggagna ttcgattcag gccg 284

<210> 134  
<211> 429  
<212> nucleic acid  
<213> Zea mays

<400> 134

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acttaccaag cgccccgagg agattggagc tggtgaccag gggcacatgt ttgggtatgc 180  
gactgacgag acccctgagc tgatgccct cagccatgtn cttgccacca agcttgggtgc 240  
tcgtctnaca aangntcgca agaaatggaa cctggcccct ggcttaagcc cgatnggnaa 300  
gaccaagtgc acaagtggaa tanccgnaac caaggggtggc nccatgggtt cccattcgtg 360  
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anttggcnt 429

<210> 135  
<211> 282  
<212> nucleic acid  
<213> Zea mays

<400> 135

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ggcggctggg gagcccacgg cgggtggcgcn ttctccggca aggacccac caaggtggac 120  
cgcagcggcg cntacgtggc caggcaggcc gccaaagca tcgtngccag cngctcgenc 180  
gcogntgcnt ngtgcaggtg tcgtacgcca tcggctgccg gagcccctgt ccgtgttngt 240  
caactcgtac ggcngcggca cgntccccga caaggagntc tc 282

<210> 136  
<211> 279  
<212> nucleic acid  
<213> Zea mays

<400> 136

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gacacctgcc gcgagatcgg gttcacctcc gacgacgtgg gcctcgacgc cgaccgctgc 120  
aaggtgctgg tgaacatcga gcagcagtc cccgacatcg cgcagggcgt gcacgggcat 180  
tcacgaagcg gcccgaggag atcggcgcgg gcgaccaggg ccacatgttc gggtagccca 240  
ccgacgagac ccccgagtga tgccgtnagc natgtngc 279

<210> 137  
<211> 283  
<212> nucleic acid  
<213> Zea mays

<400> 137

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ggcgggttggg gagccacagg cgggtggcgn ttctccggca aggacccac caaggtggac 180  
cgcagcggcg cctacgtggc caggcaggcc gccaaagca tcgtggccag cggcttcgcc 240  
cgcnctgcc tcgtgcaggt gtcgtacgcc atcgggtgcc gga 283

<210> 138

<211> 297

<212> nucleic acid

<213> Zea mays

<400> 138

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cgctggcctc actggccgga agatcatcat cgacacctac ggtggctggg gagccacagg 180  
cgggggcgcc ttctccggca aggacccgac caaggtggac cgcagcgggg cctacgtcgc 240  
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<210> 139

<211> 317

<212> nucleic acid

<213> Zea mays

<400> 139

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caagctgtgc gaccaggtgt cggacgccgt gcttgacgca tgcctcgcgc aggacccgca 180  
cagcaagggtg gcctgcgaga cctgcaccaa gaccaacatg gtgatgggtgt tcggcgagat 240  
cacgaccaag gcgaccgtgg actacgagaa gatcgtgcgc gacacctgcc gcgagatcgg 300  
gttcacctcc gacgacg 317

<210> 140

<211> 277

<212> nucleic acid

<213> Zea mays

<400> 140

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cttcgacttc aggcccgga tgatcagcat caacctcgac ctgaagaagg gcggcaacag 180  
gttcatcaag accgcccctt acggccactt cggccgtgac gacggcgact tcacctggga 240  
ggtggtgaag cccctcaagt tcgacaaggc atcgctt 277

<210> 141

<211> 279

<212> nucleic acid

<213> Zea mays

<400> 141

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acctcaagaa aggcggcaac gggcgctacc tcaagacggc ggccctacggc cactttggaa 180  
gggacgaccc tgacttcacc tgggaggtgg tgaagccact caagtggag aaaccttctg 240  
cctaaggcgg ctttttttcc agtaagaagc ttttgggtg 279

<210> 142

<211> 263

<212> nucleic acid

<213> Zea mays

<400> 142

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ctccgtgttc gtcgacacgt acggcaccgg cgcgatcccc gacaaggaga tcctcaagat 180  
tgtcaaggag aacttcgatt tcaggcctgg catgatcatc atcaaccttg acctcaagaa 240  
aggcggcaac gggcgctacc tca 263

<210> 143

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 143

cgatcatcaag ccagtcaccc ccgagcagta cctcgacgag aagacaatct tccacctcaa 60  
 cccgtctggc cgtttcgtca tcggcgagacc tcacggcgac gccggcctca ctggccggaa 120  
 gatcatcatc gacacctacg gtggctgggg agcccacggc gggggcgctt tctccggcaa 180  
 ggacccgacc aaggtggacc gcagcggggc ctacgtcgcg aggcaggctg ccaagagcat 240  
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<210> 144

<211> 280

<212> nucleic acid

<213> Zea mays

<400> 144

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 ggtgctcgtc tcaccgaggt ccgcaagaac ggaacctgcc cctggctcag gcctgatggg 180  
 aagaccaggt tgacagtcca gtaccgcaat gaggggtggg ccatgggtccc catccgtgtc 240  
 cacaccgtcc tcattctcac ccgcacgacg agacagtgac 280

<210> 145

<211> 251

<212> nucleic acid

<213> Zea mays

<400> 145

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 atcggcgcggt ggcaccaggg ccacatgttc gggtagccca ccgacgagac ccccgagctg 180  
 atgccgctga gccacgtggt ggccaccaag ctggggcgcg gcctcaccga ggtgcgcaag 240  
 aacggcactg g 251

<210> 146

<211> 270

<212> nucleic acid



<213> Zea mays

<400> 146

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aatgatgaga tccgtgtctga cctgaaggag catgtcatca agcctgtcat ccctgagcag 120  
taccttgacg agaagaccat cttccacctt aacctcatctg gccgttttgt cattgggtgga 180  
cctcacggcg atgctggcct cactggccgc aagatcatca ttgacaccta cgggtggctgg 240  
ggagcccatg gtggtggcgt ttctccggca 270

<210> 147

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 147

agacctgccg caacattggt tttgtgtcaa acgatgttg gcttgacgcc gaccactgca 60  
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tcaccaagcg ccccgaggag attggagctg gtgaccagg acacatgttc gggatatgca 180  
ccgatgagac ccctgagttg atgcccctca gccatgtcct tgccaccaag ctaggtgctc 240  
gtctcaccga ggtccgcaag aacggaactg cccctggctc agcctgatgg gaagaccagt 300  
gacagtcgag 310

<210> 148

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 148

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caccggcgcg atccccgaca aggagatcct gaagatcgtg aaggagaact togacttcag 120  
gcccggcatg atcatcatca acctcgacct caagaaaggc ggcaacgggc gctacctcaa 180  
gacggcgggc tacgggcact ttgggaggga cgaccccgac ttcacctggg aggtgggtgaa 240  
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<210> 149

<211> 279  
 <212> nucleic acid  
 <213> Zea mays  
  
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 aaggagcatg tcataagcc tgtcatcctg agcagtacct tgacgagaag accatcttcc 180  
 accttaacnc atctggccgc tttgtcattg gtggacctca cggcgatgct ggccctcactg 240  
 gccgcaagat catcattgac acctacggtg gctggggag 279

<210> 150  
 <211> 322  
 <212> nucleic acid  
 <213> Zea mays  
  
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 aggcaatggc ggccgagagc ntctttttca cctcggagtc cgtgaacgag gggcaccctg 120  
 acaagctgtg cgaccaggtg tcggacgccg tgcttgacgc atgcctcgcg caggaccctg 180  
 acagcaaggt ggctgcgag acctgcacca agaccaacat ggtgatgggtg ttcggcgaga 240  
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 ggttcactcc gacgacgtgg gc 322

<210> 151  
 <211> 283  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 151  
  
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 gcttcctttt cacctcggag tccgtgaacg aggggcaccc cgacaagctg tgcgaccagg 120  
 tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc cgacagcaag gtggcctgcg 180  
 agacctgcac caagaccaac atggtgatgg tggtcggcga gatcacgacc aaggcgaccg 240  
 tggactacga gaagatcgtg cgcgacacct gccgagat cg 283

<210> 152  
 <211> 316  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 152  
  
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 aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180  
 cgcaggaccc cgacagcaag gtggcctgcg agacctgcac caagaccaac atggtgatgg 240  
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 gccgcgagat cggggt 316

<210> 153  
 <211> 277  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 153  
  
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 agctaggtgc tcgtctcacc gaggtccgca agaacggaac ntgcccctgg ctcagggctg 180  
 atgggaagac ccaggtgaca gtcgagtacc gcaatgaggg tggtgccatg gtcccatcc 240  
 gtgtccacac cgtcctcatc tccaccagc acgacga 277

<210> 154  
 <211> 272  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 154  
  
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 gatcatcatt gacacctacg gtggctgggg agcccatggg ggtggcgctt tctccggcaa 180  
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gtcgccagcg gccttgctcg ccgcgccatc gt

272

<210> 155  
<211> 297  
<212> nucleic acid  
<213> Zea mays

<400> 155

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aagctgtgcg accaggtgtc ggacnccgtg cttgacgcat gcctcgcgca ggaccccgac 180  
agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg tgatggtggt cggcgagatc 240  
acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acacctgccg cgagatc 297

<210> 156  
<211> 267  
<212> nucleic acid  
<213> Zea mays

<400> 156

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gcctcgcgca ggaccccgac agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg 180  
tgatggtggt cggcgagatc acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg 240  
acacctgccg cgagatcggg ttacact 267

<210> 157  
<211> 261  
<212> nucleic acid  
<213> Zea mays

<400> 157

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gcaagaatgg aacctgcccc tggctcaggc ccgatgggaa gaccaggtg acagtggagt 180  
accgcaacga ggggtggcgcc atgggttccca tccgtgtgca cacagtctc atctctaccc 240

agcacgacga gacagtcacc a

261

<210> 158  
<211> 288  
<212> nucleic acid  
<213> Zea mays  
  
<400> 158

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cggaacctgc ccctggctca ggctgatgg gaagaccag gtgacagtcg agtaccgcaa 120  
tgaggggtgg gccatggtcc ccacccgtgt ccacaccgtc ctcactcca cccagcacga 180  
cgagacagtg accaatgatg agatcgctgc tgacctgaag gagcatgtca tcaagcctgt 240  
catccctgag cagtacttga cgagaagaca tttccactt aaacccat 288

CCAGGAGGCAATGGCGGCCGAGAGCTTCTTTTCACNTCGGATCCCGTGGAACGAGGGGCACCCGACACAAGCTGTGCGACCAGGTGTCGGACGCCGTGTTGACGCGATGCCTCGCGCAGGACCCGACAGCAAGGTGGCAGACNTGCAACAAGACCAACNTGGTGATGTGTTTCGGCGAGATCACGACCAAGGCGACGTGGACTACGAGAAGATCGTGCAGGACACCTGCCGCGAGATCGGGTTTCACT

<210> 159  
<211> 311  
<212> nucleic acid  
<213> Zea mays  
  
<400> 159

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cgcacaagct gtgcgaccag gtgtcggacg ccgtgcttga cgcagtcctc ggcgaggacc 180  
ccgacagcaa ggtggcntgc gagacntgca ccaagaccaa cntggtgatg gtgttcggcg 240  
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<210> 160  
<211> 267  
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<213> Zea mays  
  
<400> 160

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gtacggcacc ggcacgatcc ccgacaagga gatcctcaag atcgtgaagg agaacttcga 240  
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<210> 161  
<211> 284  
<212> nucleic acid  
<213> Zea mays  
  
<400> 161

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ccaaggccaa tgttgactac gagaagattg tcagggagac ctgccgcaac attggttttg 240  
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<210> 162  
<211> 237  
<212> nucleic acid  
<213> Zea mays  
  
<400> 162

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gcgccttctc cggcaaggac cccaccaagg tggaccgcag cggcgcttac gtggccaggc 180  
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<210> 163  
<211> 236  
<212> nucleic acid  
<213> Zea mays  
  
<400> 163

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accgtcacca acgacgagat cgccgccgac ctcaaggagc acgtcatcaa gcccgatgc 180  
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<210> 164  
 <211> 272  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 164  
  
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<210> 165  
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 <213> Zea mays  
  
 <400> 165  
  
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<210> 166  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 166  
  
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 gacagcaagg tggcctgcga gacctgcacc aagaccaaca tggatgatgg gttcggcgag 240  
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<210> 167  
 <211> 298  
 <212> nucleic acid  
 <213> Zea mays  
  
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 ccgacagcaa ggtggcctgc gagacctgca ccaagaccaa catggtgatg gtgttcggcg 240  
 agatcacgac caaggcgacc gtggactacg agaagatcnt gcgcgacacc tgccgcga 298

<210> 168  
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 <213> Zea mays  
  
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 cagtcgagta ccgcaatgag ggtgggtgcca tggccccat ccgtgtccac accgtcctca 120  
 tctccacca gcacgacgag acagtgacca atgatgagat cgctgctgac ctgaaggagc 180  
 atgtcatcaa gcctgtcatc cctgagcagt accttgacga gaagaccatc ttccacctta 240  
 acccatctgg ccgtttgtca ttggt 265

<210> 169  
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 <213> Zea mays  
  
 <400> 169  
  
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 cccgacagca aggtggcctg cgagacctgc accaagacca acatggtgat ggtgttcggc 120  
 gagatcacga ccaaggcgac cgtggactac gagaagatcg tgcgcgacac ctgccgcgag 180  
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 catcgagcag c 251





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250

<210> 173  
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<212> nucleic acid  
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<400> 173

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cgagaagaag gcaatggcgg cggagagctt cctgttcacc tcggagtccg tgaacgaggg 120  
gcacccagac aagctgtgcg accaggtgtc ggacgcggtg ctggacgcct gcctggcgca 180  
ggaccccgac agcaaggtgg cctgcgagac ctgcaccaag acgaacatgg tgatggtgtt 240  
cggcgagatc accaccaagg cgagcgtgga ctacgagaag atcgtgcgcg acacctgccg 300  
cgag 304

<210> 174  
<211> 328  
<212> nucleic acid  
<213> Zea mays

<400> 174

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cggtagagcg agaagaaggc aatggcgggc gagagcttcc ttttcacctc ggagtccgtg 120  
aacgangggc accccgacaa gctgtgcgaa ccaggtgtcg gacgccgtgc ttgacgcatg 180  
cctgcgcgag gaccccgaca gcaangtggc ctgcgagacc tgcaccaaga ccaacatggt 240  
gatggtgttc ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga 300  
cacctgccgc gagatcgggt tcactccg 328

<210> 175  
<211> 297  
<212> nucleic acid  
<213> Zea mays

<400> 175

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gaacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg cgcaggaccc 180  
cgacagcaag gtggcctgcg agacctgcac caagaccaac atgggtgatgg tgttcggcga 240  
gatcacgacc aaggcgaccg tggactacga gaagatcgtg cgcgacacct gccgcga 297

<210> 176  
<211> 275  
<212> nucleic acid  
<213> Zea mays  
<400> 176

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ctacgagaag attgtcaggg agacctgccg caacattggt tttgtgtcaa acgatgttgg 180  
gcttgacgcc gaccactgca aggtgctcgt gaacattgag cagcagtccc ctgatatgct 240  
caggggtgtgc atggccattc accaagcgcc ccgag 275

<210> 177  
<211> 534  
<212> nucleic acid  
<213> Zea mays  
<400> 177

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gagaagaagg caatggcgcc cgagagcttc cttttcacct cggagtccgt gaacgagggg 180  
caccocgaca agctgtgcga ccaggtgtcg gacgccgtgc ttgacgcatg cctcgcgcag 240  
gaccccgaca gcaaggtggc ctgcgagacc tgcaccaaga ccaacatggt gatggtgttc 300  
ggcgagatca cgaccaaggc gaccgtggac tacgagaaga tcgtgcgcga caccttnccg 360  
cgagatcggg ttcaccttcc gacgacntgg gccttgacgc ccaccgggtt caaggtgctt 420  
gtgnacattg agcaagaatt ccccgaaatt gngcaaggcg ttcaccggca ctttacgaac 480  
cggcccnagg aagatcggnc cggccnacca nggncaattt tttgggtccc cccc 534

<210> 178  
<211> 248  
<212> nucleic acid

<213> Zea mays

<400> 178

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gaatggaacc tgcccctggc tcaggcccga tgggaagacc caggtgacag tggagtaccg 120  
caacgagggg ggcccatgg ttcccatccg tgtgcacaca gtcctcatct ctaccagca 180  
cgacgagaca gtcaccaacg acgagattgc tgetgacctg aaggagcacg tcatcaagcc 240  
agtcatcc 248

<210> 179

<211> 302

<212> nucleic acid

<213> Zea mays

<400> 179

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gnaccagac aagctgtgcg accaggtgtc ggacgcggtg ctggacgcct gcctggcgca 180  
ggaccccgac agcaaggtgg cctgcgagac ctgcaccaag acgaacatgg tgatggtgtt 240  
cggcgagatc accaccaagg cgagcgtgga ctacgagaag atcgtgcgcg acacctgccg 300  
cg 302

<210> 180

<211> 281

<212> nucleic acid

<213> Zea mays

<400> 180

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cccggcatga tcatcatcaa cctcgacctc aagaaaggcg gcaacgggcg ctacctcaag 180  
acggcggcct acgggcactt tgggaggagc gaccccgact tcacctggga aggtggtgaa 240  
gcccctcaag gcggagaagc cgtcttctgc atgaggcgcn t 281

<210> 181

<211> 269  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 181  
  
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 ccaggtctca gatgccgttc ttgacgcttg ccttgctgag gaccctgaca gcaaggttgc 120  
 ttgtgagacc tgcaccaaga ccaacatggt catggtcttt ggtgagatca ccaccaaggc 180  
 caatgttgac gccgagaaga ttgtcagggg gacctgccgc aacattgggt ttgtgtcaaa 240  
 cgatgttggg cttgacgcng accatgcaa 269

<210> 182  
 <211> 286  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 182  
  
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 aagotgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180  
 agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg tgatgggtgtt cggcgagatc 240  
 acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acacct 286

<210> 183  
 <211> 240  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 183  
  
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 gcaattcacg aagcggccccg aggagatcgg cgcgggcgac cagggccaca tgttcgggta 180  
 cgnacccgac gagacccccg agctgatgcc gctgagccac gtggtggcca ccaagctggg 240

<210> 184  
 <211> 250  
 <212> nucleic acid

<213> Zea mays

<400> 184

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ngtggctggg gagcccatgg tgggtggcgt ttctccggca aggacccaac caagggtgac 180  
cgcagcggag cctatgtcgc aaggcaggct gccaaagagca tcgtcgccag cggccttgct 240  
cgccgcgcca 250

<210> 185

<211> 305

<212> nucleic acid

<213> Zea mays

<400> 185

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gagcgagcga gaagaaggca atggcgggcg agagcttcct gttcacctcg gagtccgtga 120  
acgaggggca cccagacaag ctgtgcgacc aggtgtcgga cgcggtgctg gacgcctgcc 180  
tggcgcagga ccccgacagc aagggtggcct gcgagacctg caccaagacg aacatgggtga 240  
tggtgttcgg cgagatcacc accaaggcga gcgtggacta cgagaagatc gtgcgcgaca 300  
cctgc 305

<210> 186

<211> 305

<212> nucleic acid

<213> Zea mays

<400> 186

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atcgggttca cctcccgacg acgtgggcct cgacgccgac cgctgcnagg ngctgggtgaa 180  
natcgagcan cagtcncccg acatcgcgca ngcntgcacg ggcacttcac naagcgnccc 240  
gangagatcg ncgcggcncta ccatnggcac atgttcgggt acncnaccna nnagacnnnc 300  
gagct 305

<210> 187  
 <211> 274  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 187  
  
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 ggttcccata cgtgtgcaca cagtctcat ctctaccag cacgacgaga cagtcancaa 180  
 cgaagagatt gctgctgacc tgaaggagca cgtcatcaag ccagtcattc ccgagnagnn 240  
 acctcgacga gaagacaata ttccacacac ttna 274

<210> 188  
 <211> 232  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 188  
  
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 tcaaccgctc tggcgcgttc gtcctcggcg gacctcacgg cgacgctggc ctactggcc 120  
 ggaagatcat catcgacacc tacggtggct ggggagccca cggcgggggc gccttctccg 180  
 gcaaggaccc gaccaaggtg gaccgcagcg gggcctacgt cgcgaggcag gc 232

<210> 189  
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 <212> nucleic acid  
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 <400> 189  
  
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 tggaaacctgc ccttggtca ggccgatgg gaagaccag gtgacagtgg agtaccgcaa 120  
 cgagggtggc gccatggttc ccatccgtgt gcacacagtc ctcatctta ccagcacga 180  
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 cat 243

<210> 190

<211> 290  
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 <400> 190  
  
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 aagctgtgcg accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180  
 ancnaggtgg cctgcgagac ctgcaccaag accaacaatgg tgatggtggt cggcnagatc 240  
 acgaccaagg cgaccgtgga ctacnagaag atcgtgcgcg acacctgccg 290

<210> 191  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 191  
  
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 tgtgagacct gcaccaagac caacatggtc atggtctttg gtgagatcac caccaaggcc 180  
 aatgttgact acgagangat tgtcaggag acctgccgca acattggttt tgtgtcaaac 240  
 gatgttgggc tgacgccgac cactgc 266

<210> 192  
 <211> 276  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 192  
  
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 agcagtaact tgacgagaag accatcttcc acttaacca tctgcccgtt ttgtcattgg 180  
 tggacctcac ggcgatgctg gcctcactgg ccgcaagatc atcattgaca ctacggtggc 240  
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<210> 193



<211> 292  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 193  
  
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 ttctgctgaca cgtacggcac cggcgcgatc cccgacaagg agatcctcaa gattgtcaag 240  
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<210> 194  
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 gcagggcgctg caggggcact tcacgaagcg gcccagaggag atcggcgcgcg gcgaccaggg 180  
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<210> 195  
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 ccgacaagct gtgcgaccag gtgtcgagcg ccgtgcttga cgcattgctc gngcaggacc 180  
 ccgacagcaa ggtggcctgc gagacctgca ccaagaccaa catggtgatt gtgttcggcg 240  
 agatcacgac canggcgacc gtggactacg agaagatcgt gcgcnacac 289

<210> 196  
 <211> 300  
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<213> Zea mays

<400> 196

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aacgaggggc acccagacaa gctgtgcgac caggtgtcgg acgcggtgct ggacgcctgc 180  
ctggcgagc accccgacag caaggtggcc tgcgagacct gcaccaagac gaacatggtg 240  
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<210> 197

<211> 284

<212> nucleic acid

<213> Zea mays

<400> 197

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aagctgtgct accaggtgtc ggacgccgtn cttgangcat gcctcgcgca ggaccccgac 180  
agcaaggtgg cctgcgagac ctgnaccaag acnaacatgg tgatggtggt cggcgagatc 240  
acgaccaagg cgaccgtgga ctacgagaag atcgtgcgcg acac 284

<210> 198

<211> 282

<212> nucleic acid

<213> Zea mays

<400> 198

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accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac agcaaggtgg 180  
cctgcgagac ctgcaccaag accaacaatng tgatggtggt cggcgagatc acgaccaang 240  
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<210> 199

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 199

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cagcatcaac ctgcacctga agaagggcgg caacaggttc atcaagaccg nccctacgg 180  
ccacttcggc cgtgacgacg ccgacttcac ctgggagggtg gtgaagcccc tcaagttcga 240  
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<210> 200

<211> 291

<212> nucleic acid

<213> Zea mays

<400> 200

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angggcaccg cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180  
cgcaggaccg cgacagcaag gtggcctgcg agacctgcac caagaccaac atggtgatgg 240  
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<210> 201

<211> 337

<212> nucleic acid

<213> Zea mays

<400> 201

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catgatcatc atcaaccttg acctcaagan aggcggnaac gggcgctacc tcaagacggc 180  
ggattanggc cactttggaa gggangaccg tgacttcacc tgggatgtgg tnaagccact 240  
caantcggag aaacctnctg cctaaggcgg nttnttttc agtaagaagc ttttggtggt 300  
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 ccgacaagct gtgcgaccag gtgtcggacg ccgtgcttga cgcattgcctc gcgcaggacc 180  
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 agatcacgac caaggcgacc gtggactacg agaagatcg 279

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ggancccgac agcaaggtgg cctgcgagac ctgcaccaag accaacaatgg tgatggtggt 240  
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<210> 205  
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 <400> 205

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 ctggcgagcagg accccgacag caaggtggcc tgcgagaccn gcaccaagac gaacatggtg 240  
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 <400> 206

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 gcgcaggacc ccgacagcaa ggtggcctgc gagacctgca ccaagacgaa catggtgatg 240  
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tgagcagcag tcccctgata ttgctcaggg tgtgcatggc cacttcacca agcgccccga 180  
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ttgatgc 247

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tgacagcaag gttgcttgcg agacctgcac caagaccaac atggtcatgg tctttggtga 240

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 accctgacag caaggttgct tgtgagacct gnaccaagac caacatggtc atggtctttg 240

gtgagatcac caccaaggcc attgtgacta cgagaagatt gtccaggag acttgccgca 300  
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<400> 246

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agngtgtgca cggccacttc accaagcgcc ccgaggagat tggagctggt gaccaggggc 180  
acatgttttg gtatgcnact gacgagaccc ctgagctgat gcccctcagc catgtccttg 240  
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aggcgaccgt ggactacgag aagatcgtgc gcgacacctg ccgagagatc gggttcacct 180  
ccgacgacgt gggctcgacg ccgaccgctg c 211

<210> 248  
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<212> nucleic acid  
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<400> 248

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cactttggcc gtgacgacgc cgacttcacc tgggagngg tcaagcccct aaagaaggca 240

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<210> 249  
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<400> 249

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gacggcgggc tacgggcact ttgggaggga cgaccccgac ttcacctggg aggtgggtgaa 180  
gcccctcaag gcggagaagc cgtcttctgc atgaggcgcc tcctctgttt tggaagaagc 240  
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gcgcgctcac cgagncacgc aagaacggca ctgcgcgtgg ctnaggccnc gacggcaaga 180  
cccaggtgac ggtggagtac gtgaacgatg gcggcgccat ggtgcccgtc cgcgtcgaca 240  
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<210> 251  
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cggtgttcgt ggacacgtac ggcaccggcg cgatccccga caaggagatc ctgaagatcg 180  
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cggcaacggc acgattcccc acaagggaga ttctcaaaga tcgttaagga gaaactcgaa 300  
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aaggttnaat caaaaancgc ccgcctaagg ncaattnggc ccgttaacca acgcganttc 420  
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ccggcctcac cggccgcaag atcatcatcg acacctacgg cggctgggga gccacggcg 180  
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<210> 254  
<211> 295  
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<213> Zea mays  
  
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 cgggtgctgga cgctgcctg ggcgaggacc ccgacagcaa ggtggcctgc gagacctgca 240  
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<400> 255

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 gaccncacgg cgatgcnggc ctacnnggc gcaagancan catngacacc nacggtggcn 180  
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<210> 256  
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<400> 256

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 aacatcgagc agcagtcacc cgacatcgcg cagggcgctgc acgggcactt cacgaagcgg 180  
 cccgaggaga tcggcgcggg cgacca 206

<210> 257  
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<400> 257

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 ggcaggctgc caagagcatc gtcgccgc 208

<210> 258  
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 <212> nucleic acid  
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 <400> 258

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 gtcatccccg agcagtacct cgacgagaag acaatcttcc acctcaaccg gtctggccgt 180  
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aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc cgtgcttgac gcatgcctcg 180  
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 atttcaggcc tggcatgac atcatcaacc ttgacctcaa gaaaggcggc aacggggcgct 240  
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 gaagccattc aatcggagaa actttgctaa gcggc 335

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 tctgtgacca ggtctcagat gccgttcttg acgcttgcc tcttgaggac cctgacagca 180  
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caccctgaca agctctgtga ccaggtctca gatgccgttc ttgacgcttg ccttgctgag 180  
gaccctgaca gcaaggttgc ttgtgagacc tgcaccaaga ccaacatggt catggtcttt 240  
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<213> Zea mays  
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cattgtccag gtctcctacg ccatcggcgt gcccgagccc ctttcggtgt tcgtggacac 180  
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 ggggtgctgg acgcctgcct ggcgaggac cccgacagca aggtggcctg cgagacctgc 180  
 accaagacga acntngtgat ggtgttcggc gagatcncca ccaaggcgag cgtg 234

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 caacattgga cttggtggtg atctcaccaa agaccatgac catgttggtc ttggtgcagg 180  
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 aagctgtgag accaggtgtc ggacgccgtg cttgacgcat gcctcgcgca ggaccccgac 180  
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<400> 275

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<210> 276

<211> 228

<212> nucleic acid

<213> Zea mays

<400> 276

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cctatgtcgc aaggcaggct gccaaagagca tcgtcgccag cggccttgct cgcgcgcgca 180  
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<210> 277

<211> 253

<212> nucleic acid

<213> Zea mays

<400> 277

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cgatgctggc ctactggcc gcaagatcat cattgacacc nacggtggct ggggatcccn 180  
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<211> 268

<212> nucleic acid

<213> Zea mays

<400> 278

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 ccgtgaacga ggggcaccca gacaagctgt gcgaccaggt gtcggacgcg gtgctggacg 180  
 cctgcctggc gcaggacccc gacagcaagg tggcctgcga gacctgcacc aagacgaaca 240  
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<400> 279

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 ggggecttgc tcgccgcgcc atcgccagn tgtcttacgc canngggtgc nngancctct 240  
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 acaagctgtg cgaccaggtg tcggacgccg tgcttnangc atgcctcgcg caggacccccg 180  
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 cgagggggcac ccagacaagc tgtgcgacca ggtgtcggac gcggtgctgg acgcctgcct 180  
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cccgctctggc cgcttcgtca tcngcggacc tcacggcgac gctggcnctnn ctgnnnccggn 180  
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 acttcacctg ggaggtggtg aagccacttc aagtcggaga aaccttctgc ctaaggcggc 240  
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<400> 293

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 aacgaggggc acccagacaa gctgtgcgac cagggtgcgg acgcggtgct ggacgcctgc 180  
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 gggagacatg ccgcaaaatt ggtttcngt naganngatg tggggcttga ngntgaccac 240  
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<212> nucleic acid

<213> Zea mays

<400> 302

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tganaatngn anaagganaa nttnanantn caggcccggt tgatcattat naacctagac 180

ctcaanaaag gcggaaacgg gcnctaccta aagacggggg tctacgggcn ctttgngagg 240

gacgaccgag anttcacctg agaggtggna aagcccctca aggcggaaaa gccgtcttct 300

gcattgaggc cctcctctgt ttengaagaa gcttttggtc tggctctgct gcgctctatc 360

atgctttttt atggctncta cgtgttggtga ttcttgatct gcccttgct tatcatttgt 420

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<211> 297

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<213> Zea mays

<400> 303

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natngnceca ggtgtattat cgccatccgg cnngncngng cntctcnccg tgttcgtcga 120

caagtaacggc accggcgcca tccccgacaa gngnncnct gaagattgta caaggagaac 180

ttcgatttca ggctggcat gatcatnnnc aaccttganc tcaagaaagg nggcaacggg 240

cgctacctca agacggcggc ctacggcnac tttggaaggg acgaccctga cttcacc 297

<210> 304

<211> 259

<212> nucleic acid

<213> Zea mays

<400> 304

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tgatgggaag acccaggtga cagtcgagta ccgcaatgag ggtggtgcca tgggtcccat 180

ccgtgtccac accgtctca tctccacca gcacgacgag acagtgacca atgatgagat 240  
cgctgctgac ctgaaggag 259

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<213> Zea mays  
  
<400> 305

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gttc 244

<210> 306  
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<213> Zea mays  
  
<400> 306

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acaccctgac aagctctgag accaggtctc agatgctgtt ctggacgctt gccttgctga 180  
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<210> 307  
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cctcacggcg aggtgggcnt nactggccgg angatntcat cganannagg tgtttgggga 300  
nccacggggg 310

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<212> nucleic acid  
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<400> 309

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ccccgacagc aaggtggcct gcgagacctg caccaagacc aacatgggtga t 231





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gtntncatgg ccacttcacc aagcgccccg aggagattgg agctggngac cagggacaca 180  
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269

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<400> 317

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<400> 318

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<400> 319

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<210> 330

<211> 176

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<213> Zea mays

<400> 330

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gtgcgcaaga acggcacctg cgcttggtg aggcccgacg gcaagacca ggtgac 176

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<212> nucleic acid

<213> Zea mays

<400> 331

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ccnganaagc tgtgcncca ngntcggac gccgtgcttg acgcatgcct cgcgcaggan 180

cccacagca aggtggatgc gagacctgca taagaccaac atggtgatgg tgttcgncga 240

gatcacgacc aaggcgncg tgg 263

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<211> 225

<212> nucleic acid

<213> Zea mays

<400> 332

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aggggcaccc cgacaagctg tgcgaccagg tgcggaagc cgtgcntgac gcatgcctcg 180

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 gaagatggcc ggactcgaca ccttcctctt cacctcggag tccgtgaacg agggacaccc 240  
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<400> 337

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<400> 338

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gcagatcaaa gaagatggca gctgtcgaca cattcctctt cacctcggag tctgtgaacg 180  
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 tgggtgaagcc cctcaagttc gacaaggcat cggcttaagg ttgggagtgt cactgtggac 180  
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<213> Zea mays  
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143

<210> 346

<211> 142

<212> nucleic acid

<213> Zea mays

<400> 346

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acctcaagaa aggcggcaac gg 142

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<211> 288

<212> nucleic acid

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gcagatcaaa gaagatggca gctgtcgaca cattcctctt cacntcggag tctgtgaacg 180

agggacacnc tgacaagctc tgtgaanagg tctcagatgc cgttcttgac gcttgcnttg 240

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<211> 288

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<400> 348

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cgccgccgcc cagatcaaag aagatggcag ctgtcgacac attcctcttc acctcggagt 180

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aagccatgtc gcnccactga ccggcttaat gattggtata atttgggtgtg gcaacancca 420  
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<213> Zea mays

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tcaagacggg ggcctacggg cactttggga gggacgaccc cgacttcacc tgggaggtgg 180

tgaagcccct caaggcggag aagccgtctt ctgcatgagg cgctcctct gtttcggaag 240

aagcttttgg tctggtctgc ctgcgctcta tcatgctttt ttatggtccc tacgtgttgt 300

gattcttgat ctgccccttg cttatcattt gtattgtact gtcactgtcc taataagtgg 360

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gaccaccccg cgccgccgca gatcaaagaa gatggcagct gtcgacacat tctctttcac 180

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<213> Zea mays

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aagttcgaca aggcacggc ttaaggttgg gagtgtcact gtggacatga ggactacctt 180

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 <400> 369  
  
 gacgagacga gtcccctnac cccacctac gcctcaccca accggaacga acaagttaca 60  
 atactgcac ccaacccgc ctcgaccgga tctcgtcgga ctcggatccg cccgaccacc 120  
 ccgcgcgcc gcagatcaaa gaagatggca gctgtcgaca cattctctt cacctcggag 180  
 tctgtgaacg agggacaccc tgacaagctc tgtgaccagg tctcagatgc cgttcttgac 240  
 gcttgccctg ctgaggaacc tgacagcaag gttg 274

<210> 370  
 <211> 203  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 370  
  
 ctctttctccc tcttgccgnt cccgaataaa gagcagcagc gcaagaggtc ggtagagcga 60  
 gaagaaggca atggcggcgc agagcttcct tttcaentcg ggtccgtga acgaggggca 120  
 cgccgacaag ctgtgcgacc aggtgtcgga cgccgtgctt gacgcatgcc tcgcgagga 180  
 cnccgacagc aagtggcctg cga 203

<210> 371  
 <211> 201  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 371  
  
 gcctcttctc ctccctcctg ccgggtcctt aataaagagc agcagcgcaa gaggttggtgta 60  
 gagcgagcga gaagaaggca atggcggcgc agagcttcct gttcacctcg ggtccgtga 120  
 acgaggggca cccagacaag ctgtgcgacc aggtgtcgga cgcggtgctg gacgcctgcc 180  
 tggcgagga ccccgacagc a 201

<210> 372  
 <211> 307  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 372  
  
 tcagcatcaa cctcgacctg aagaagggcg gcaacaggtt catcaagacc gccgcctacg 60  
 gccacttcgg ccgtgacgac gccgacttca cctgggaggt ggtgaagccc ctcaagttcg 120  
 acaaggcatc ggcttaaggt tgggagtgtc actgtggaca tgaggactac cttcctctgg 180  
 ctctgctgtt acctgcaagc attgctgctg ctggatgtgt gtgtttgatc agtgactggc 240  
 tgctgctcca tagaagatga acggagagaa ggatgatgaa gggctttggc aatcgccgct 300  
 gcaactg 307

<210> 373

<211> 283  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 373  
  
 gtcttaacgcc atcgggtgcc cgagcctctc tccgtgttcg tcgacacgta cggcaccggc 60  
 gogatccccg acaaggagat nctcaagatt gtcaagngaa ttcgatttca ggcttggcat 120  
 gatcatcatc aaccttgact caagaaaggc ggcaacgggc gctactcaag acgcggccta 180  
 cggcactttg gaaggagacc tgattcactg ggagtgtga accatcaatg gagaacttgc 240  
 tcnnngctat tatataantt gtgcgttctg actatngtta nag 283

<210> 374  
 <211> 181  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 374  
  
 ctgtgccggtc ccgaataaaag agcanncagc gcaagagntc ggtagagcga caagaaggca 60  
 atggcgggcg agagcttctt ttccacctcg gactccgtga acgaggggca ccccgacaag 120  
 ctgtgcgacc aggtgtcgga cgccgtgctt gacgcagtcg tcgcgcagga ccccgacagc 180  
 a 181

<210> 375  
 <211> 201  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 375  
  
 ctgccgggtc cttaataaaag agcagcagcg caagngagc cgccagcttg cccaggttg 60  
 gtagagcgag cgagaagaag gcaatggcgg cggagagctt cctgttcacc tcggagtccg 120  
 tgaacgaggg gcacccagac aagctgtgcg accaggtgtc ggacgcggtg ctggacgcct 180  
 gcctggcgca ggaccccgac a 201

<210> 376  
 <211> 216  
 <212> nucleic acid  
 <213> Zea mays

<400> 376

acatccatcc ccgttcgcct cttctcctcc ctctgcccgg gtccttaata aagagcagca 60  
gcgcaagagg ttggtagagc gagcgagaag aaggcaatgg cggcggagag cttcctgttc 120  
acctcggagt ccgtgaacga gtggcaccca gacaagctgt gcgaccaggt gtcggacgcg 180  
gtgctggacg cctgcctggc gcaggacccc gacagc 216

<210> 377

<211> 130

<212> nucleic acid

<213> Zea mays

<400> 377

gtcgaacgat gtcgggcttg acgctgacca ctgcaagggtg cntgtgaaca ttgagcagca 60  
gtccccgat attgcnacagg gtgtgcacgg ccacttcacc aagcgccccg aggagattng 120  
agctggtgac 130

<210> 378

<211> 306

<212> nucleic acid

<213> Zea mays

<400> 378

gtcggatctg agacgagacg agacgatccc ccctcncctc aaccggaact tgttttacct 60  
catctcatcc cactgantcc acccaccac ccgcccgtg cntccgccg atntcgtcgg 120  
actcggatcc gcccgaccac gaccaccccg cgtcgccgcc gcgcanagca gcagatcaga 180  
gaagatggcc ggactcgaca cttctctctt cacctcggag tccgtgaacg agggacaccc 240  
tgacaagctc tgcgaccagg tctcagatgc tgttctggag gnttgccttg ctgaggaccc 300  
tgacag 306

<210> 379

<211> 313

<212> nucleic acid

<213> Zea mays

<400> 379

gagacgagac gannnnccct cccctcaacc ggaacttgtt ttaccccatc tcatccact 60

gactccan cn acccaccgc ncgtgcctc cgccggatct cgtcggactc ggatccgccc 120  
gaccacgacc accccgcgtc gccgccgcgc agagcagcag atcagagaag atggccggac 180  
tcgacacctt cctcttcacc tcggagtccg tgaacgaggg acaccctgac aagctctgga 240  
ccaggtctca gatgctgttc tggacgcttg ccttgctgag gacctgacag caaggttgct 300  
tgggagacct gca 313

<210> 380  
<211> 134  
<212> nucleic acid  
<213> Zea mays  
<400> 380

gengacctca nggagcacgt catcaagcnc gtgatccctg agaagtacct cgacgagang 60  
accatcttcc acctcaacct gtccggggcgc ttctgcatcg gcngggcccca cggtnacncc 120  
ngentcacng gtgc 134

<210> 381  
<211> 294  
<212> nucleic acid  
<213> Zea mays  
<400> 381

ccgggtcgga ttgagacga gacgagttac catctcatcc caactccgga acgaacaagt 60  
taccatctca tcccaactcc gcctcgaccg gatctcgctg gactcggatc cggccgacca 120  
ccccgcgccg ccgcagatca aagaagatgg cagctgtcac acattcctct tcacctcgga 180  
gtctgtgaac gagggacacc ctgacaagct ctgtgaccag gtctcagatg ccgttcttga 240  
cgcttgctt gctgaggacc ctgacagcaa ggttgcttgt gagactgcac caag 294

<210> 382  
<211> 164  
<212> nucleic acid  
<213> Zea mays  
<400> 382

cctgagttga tgccccctcag ccatgtcctt gccagcaaac taggtgctcg tntcaccgag 60  
gtccgcaaga gcggaaacct gccctggct acaggcctga tgggaagacc caggtgacag 120

togagtaccg cantgaggggt ggtgccatgg tccccatccg tntc 164

<210> 383  
 <211> 247  
 <212> nucleic acid  
 <213> Zea mays

<400> 383

cggagaaacc ttctgnctaa ggccgccgtn ttgaggtagn gnccgtngcc gactttcttg 60  
 nggnnnaggt tgatnatgat catgccaggc ctgaaatcga agttctcctt gacaatcttg 120  
 aggatctcct tgtcggngat cagngccagt gcacgtacag ngtcnacgaa cacnganaga 180  
 ngctcgggca ctccnntnnc ngtaagacan ctggacnatg gttagtgnaa gtnatnccnt 240  
 tgaanac 247

Sequence 1

<210> 384  
 <211> 207  
 <212> nucleic acid  
 <213> Zea mays

<400> 384

cacgtctaata cagacatttt actcagaagt catctttgct tgccgggtccc gaataaagag 60  
 cagcagcgca agaggtcggg agagcgagaa gaaggcaatg gcggccgaga gcttcctttt 120  
 cacctcggag tccgtgaacg aggggcaccc cgacaagctg tgcgaccagg tgtcggacgc 180  
 cgtgcttgac gcatgcctcg cgcagga 207

<210> 385  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays

<400> 385

gggtcggatc tgagacgagg cgagacgacc cccctcccc tcaaccggaa cttgttttac 60  
 cccatctcat cccacagnct ccacccannc gcccgctgcc tccgccgat ctctcggac 120  
 tcggatccgc ccgaccaccc cgcgccgccg ccgcgcagag cagcagcaga tcagagaaga 180  
 tggccggact cgacaccttc ctcttcacct cggagtccgt gaacgagggga caccctgaca 240  
 agctctgcga ccaggtctca gatgctgttc tggacgcttg ccttgctgag ga 292



<210> 386  
 <211> 142  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 386  
  
 cgcgcgcctc accgaggtgc gcaagaacgg nncacctgcg cctgggtgag gcccgaacggc 60  
 aagacccagg tgacgggtgga gtacntgaac gagggcgggcg ccatgggtgcc cgtccgcntg 120  
 cacaccgtgt catctccaca ca 142

<210> 387  
 <211> 137  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 387  
  
 gngggcgacc agggccacnt gnnccgggtac gccaccgacg agacccccga gctgatgcng 60  
 ctgagccacg ngntggccac caagntgggc gcgcgcntca ccgangngcg caagaacggc 120  
 acntggcnen tggngga 137

<210> 388  
 <211> 159  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 388  
  
 gaaaggcggc aacgggcgct acctcaagac ggcggcctac gggcactttg ggaggnacga 60  
 ccccgacttc acctggnagg tggatgaagc cctcaaggcg gagaagccgt cttctgcatg 120  
 aggcgccctc tctnttttgn aagangcttt tggtcnggt 159

<210> 389  
 <211> 268  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 389  
  
 gacgagacga gtcccctccc cccacctcgc ctacccaac cggaacgaac aagttacaat 60  
 ctcatcccaa ccccgccctcg accggatctc gtcggactcg gatccgccc accaccccgc 120

gccgccgcag atcaaagaag aggcagctgt cgacacattc ctcttcacct cggagtctgt 180  
 gaacncggga caccctgaca agctctgtga ccangtctca gatgccgttc ttgacgcttg 240  
 ccttgctgag gaccctgaca gcaaggtt 268

<210> 390  
 <211> 282  
 <212> nucleic acid  
 <213> Zea mays

<400> 390

aagaagggcg gcaacagggt catcaagacc gccgcctacg gccacttcgg ccgtgacgac 60  
 gccgacttca cctgggaggt ggtgaagccc ctcaagttcg acaaggcatc ggcttaaggt 120  
 tgggagtgtc actgtggaca tgaggactac ctctctctgg ctctgctgtt acctgcaagc 180  
 attgctgctg ctggatgtgt gtgtttgatc agtgactggc tgctgcttcc atagaagatg 240  
 aaggagagaa ggatgatgaa gggctttggc aatcgccgcg ca 282

<210> 391  
 <211> 272  
 <212> nucleic acid  
 <213> Zea mays

<400> 391

caacggggcg tacctcaaga cggcggccta cggccacttt ggaagggacg accctgactt 60  
 cacctgggag gtggtgaagc cactcaagtc ggagaaacct tctgcctaag gcggcctttt 120  
 tttttcagta agaagctttt ggtggtctgc tgtgcttaat catgctttta tatggcttct 180  
 acatgttgtg gttctttctt gatctgcacc gcgcttatcg tttgtgttgt actgccctaa 240  
 taagtgggtgc tatgaggact gtttctgggt tt 272

<210> 392  
 <211> 291  
 <212> nucleic acid  
 <213> Zea mays

<400> 392

cggatctgag acgagacgag acgannnncc ctccctcaa ccggaacttg ttttacccca 60  
 tctcatccca ctgnoctcn gccaccacc cgcncgctgc ctccgccga tctcgtcgga 120

ctcggatccg cccgaccacg accaccccg ctcgccgccc cgcagagcag cagatcagag 180  
aagatggccg gactcgacac cttcctcttc acctcggagt ccgtgaacga gggacaccct 240  
gacaagctct ggcaccaggt ctcagatgct gttctggacg cttgccttgc t 291

<210> 393  
<211> 531  
<212> nucleic acid  
<213> Zea mays

<400> 393

agnnnnnnn natntaatga atttnangaa tgctctacch gnaattcccg ggtcgaccca 60  
cgcgtccgnc cacnecgccc angagatcct caagatcgng aaggagaact tcgacttcag 120  
gcccnagnatg atcagcatca acctngacct gaanaaggnc ggcaacaggt tcatcaagac 180  
cgacgcctac agtcacttcn gncgtgacga cnccgactta cctggnaggt ggtgaacccc 240  
tcaagtccga caaggcatcg ncttaaggct gngaagtgc cactgtggac attgaggact 300  
accttactct ggctctgntg gtacctgcaa agcattggct gctgatggat gtntgngnct 360  
gatcaagnga ctggctgctg cttcatanna gatntaccg aganaaagat gatgnataaa 420  
ggcttnggca atcggcggtt canctgnaac ccatgccatt ccgcttanng aatggggata 480  
anttggcttg gaaanaanca tcattattat ggnetgaact ttcacttta c 531

<210> 394  
<211> 572  
<212> nucleic acid  
<213> Zea mays

<400> 394

gggggnnnng gnaacttcta tntcgnccg caccgtccaa aaaatcccg ggtccgaccc 60  
acgcgttccg aggcnaacttt tctcccggca aagggaccca aaccaaagg tttgaaccnc 120  
aagccgggaa ccctaatttt cgcaaaggg caanggctng cccaaagaac caatccgtcc 180  
gcccgaagccg ggccctttgc ctccgcccgc cgccaatccg ttccaangat tgtcttaacg 240  
ccaatccgng cgttnccccg aaacctctct ccgttggtcg tcgacacnta cggcaccngg 300  
cgcgatcccc gacaaggaan atnctcaaga ttgtcaagga agaacttca tttcaggcct 360  
gngcatgac atcatcaacc ttgacctcaa gaaangcggc aacggggcgc tacctcaaag 420

acggcggcct aaggccactt tgggaaaggg acnaaccctg aattcaacct ggggaaggtt 480  
gttgaaagcc aactcaaaag ttccgaaaaa aanccttctg gccnnaaggg cgggcccttt 540  
ttttcnagtt aanaaacctt ttgggggggg nc 572

<210> 395  
<211> 127  
<212> nucleic acid  
<213> Zea mays

<400> 395

cagcggcctt gctcgngcg ccatcgcca ggtgtcttac gccatcggcg tgcccgatnc 60  
ctctctccgt gttcgtcgac acgtacggca ccggcgcgat ccccgacaag gngatcctca 120  
agattgt 127

100  
90  
80  
70  
60  
50  
40  
30  
20  
10  
0

<210> 396  
<211> 294  
<212> nucleic acid  
<213> Zea mays

<400> 396

tcggnatctg agacgagacg agacgannnn ccctcccctc aaccggaact tgttttacct 60  
catctcatcc cactgactcc ntnnancnac ccgcangctg cctccgncgg atctcgtcgg 120  
actcggatcc gcccgaccac gaccaccccg cgtcgcgccg gcgcagagca gcagatcaga 180  
gaagatggcc ggactcgaca ccttcctctt cacctcggag tccgtgaacg agggacaccc 240  
tgacaagctc tgcgaccagg tctcagatgc tgttctggac gcttgcttgc tgag 294

<210> 397  
<211> 270  
<212> nucleic acid  
<213> Zea mays

<400> 397

cgantnacca tctcatcnc aactccggaa cgaacaagtt accatctcat ccanacttcc 60  
gcctcgaccg gatctngtcg gactcggatc cgcccganca ccccgnggcc gccgngatc 120  
ngagaagatg gcagctgtcg acacattcct cttnagctnc ggagtctgtg aacgagggac 180  
accctgacaa gncctgtgac caggtctcag atgccgtctt gacgcntgcn ttgctgagga 240

ccctganagc naaggtgctt gtganacctg

270

<210> 398  
<211> 284  
<212> nucleic acid  
<213> Zea mays

<400> 398

catcaacctc gacctcaaga ngggcggaac caggttcacg aagaccgccg catacggcca 60  
ctttggccgt gacgacgccg acttcacctg ggaggtggc aagcccctaa agaaggcatc 120  
cgcttaagaa tgtattggga agttcactgg acatgaggtt catcttcgtc tggctctgct 180  
gataacctga aggatnnnnn nnnnnnnnnn nnnnnnnnga tgtgtgtttg atcagtgact 240  
ggctgctctg ctccatagaa gatgaatgaa gagagagatg gtga 284

<210> 399  
<211> 297  
<212> nucleic acid  
<213> Zea mays

<400> 399

atctgagacg agacgnngnc nncctcncct caaccggaac ttgttttacc ccattctcatc 60  
ccantgante nagnnannca cncgcncgct gntccgncg gatctcttcg gactcggatc 120  
cgcccgancc cgaccanccc gcgcgcgcgc gcgcgagac agcagatcag agaagatggc 180  
cggactcgac accttctctt tcacctcgga gtccgtgaac gagggacacc ctgacaagct 240  
ctgcgaccag gtctcagatg ctgttctgga ngttgcttgc tgangacctg acagcaa 297

<210> 400  
<211> 279  
<212> nucleic acid  
<213> Zea mays

<400> 400

gtcggatctg agacgagacg agacgatnnc cctccccctc aaccggaact tgttttaacc 60  
catctcatcc cacngactcc ncccaccac ccgcccgtg cctccgccg atctcgtcgg 120  
actcggatcc gcccgaccac gaccaccccg cgtcgcgcgc gcgcagagca gcagatcaga 180  
gaagatggcc ggactcgaca ctttctctt cacctcggag tccgtgaacg agggacaccc 240

tgacaagctc tgcgaccagg tctcagatgc tggtctgga 279

<210> 401  
<211> 307  
<212> nucleic acid  
<213> Zea mays

<400> 401

cggatctgag acgagacgag acgatnnncc ctccctcaa ccggaacttg ttttaccoca 60  
tctcatccca ctgantctnc ccattccacc gcccgngcc tccgccggat ctctgtggac 120  
tcggatccgc ccgaccacga ccaccccgcg tcgcccgcgc gcagagcagc agatcagaga 180  
agatggccgg actcgacacc ttctcttca cctcggagtc cgtgaacgag ggacacctg 240  
acaagctctg cgaccaggtc tcagatgctg ttctggacgt tgcttgctga ggacctgaca 300  
gcaaggt 307

<210> 402  
<211> 291  
<212> nucleic acid  
<213> Zea mays

<400> 402

gtttgcctct tctccctctt gccgggtcccg aataaagagc agcagcgcaa gaggtacggt 60  
agagcgagaa gaaggcaatg gccgncgaga gcttctttt cacctcggag tccgtgaacg 120  
angggcacc cgcacaagctg tgcgaccagg ttacaaaaan ccgtgcttga cgcattgcctc 180  
gcgcagaccc cgacagcaag gtggcttncg agacttncac caagaccaca tgggtangttt 240  
tngnngntgg nncgncaaag nnaangngtt tnanaaaaat ntntnnancc c 291

<210> 403  
<211> 386  
<212> nucleic acid  
<213> Zea mays

<400> 403

caagaaagnc ggcaacgggc cgctacctca agacgggggc gnacggccac tttggaaggg 60  
acgacctga cttcacctgg gaggtggtga agccactcaa gtcggagaaa ctttctgcct 120  
aaggcggcct tttttttcag taagaagctt ttggtggtct gctgtgctta atcatgcttt 180

tatatggctt ctacatgttg tggttctntc ttgatctgca ccgngcttat cgnntnngtt 240  
gtactgncct aataaatnng tgcttatgan gacttgtnn tggntnnnt antanngtn 300  
naatgcttta aaacaatgan tgaattncaa gccannnttt ttttgagaag taannattat 360  
tngntaannn gntnngnntn tnnngg 386

<210> 404  
<211> 144  
<212> nucleic acid  
<213> Zea mays  
<400> 404

tccgtgttcg tcgacacgta cggcaccggc gcgatccccg acaaggagat cctcaagatt 60  
gtcaaggaga acttcgattt caggcctggc atncatcatc atcaaccttg acctcaagaa 120  
agggcggaac gggcgctacc tcaa 144

<210> 405  
<211> 293  
<212> nucleic acid  
<213> Zea mays  
<400> 405

agaacttcga cttcaggccc gggatgatca gcatcaacct cgacctgaag aagggcggca 60  
acaggttcat caagaccgcc gcctacggcc acttcggccg tgaacgacgc cgacttcacc 120  
tgggaggttg tgaagcccct caagtctgac aaggcatcgg cttaagggttg ggagtgtcac 180  
tgtggacatg aggactacct tcctctggct ctgctgttac ctgcaagcat tgctgtgtgt 240  
ggatgtgtgt gtttgatcag tgactggctg ctgtccatag aagatgaacg gag 293

<210> 406  
<211> 175  
<212> nucleic acid  
<213> Zea mays  
<400> 406

ggtcaccatc aacctcgacc tcaagaaggc cggcaacagg ttcacatcaaga ccgccgcata 60  
cggccatttg gncgtgacga cgccgacttc acctgggagg tggtaagcc cctaaagaag 120  
gcatccgctt aagaatgtat tgggaagttc actggacatg aggttcatct tcgtc 175

<210> 407  
 <211> 219  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 407  
  
 aggggtgtgca cggccacttc accaagcgcc ccgaggagat tggagctggt gaccaggggc 60  
 acatgttttg gntgcgactg acgagacccc tgagtgatgc cctcagccat gtcttgccac 120  
 caagctggtg tcgtctcacg gagtnccaag atggactgcc ctgntcagcc gtggaagacc 180  
 agtgcagtga tacgnagagg tggcatgtcc acggtnnnc 219

<210> 408  
 <211> 178  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 408  
  
 gccagggatg atcaccatca acctcgacct caagangggc ggcaacaggt tcatcaagac 60  
 cgccgcatac ggccactttg gctgaacgac gccgacttca cctgggaggt ggtcaagccc 120  
 ctaaagaagg catccgctta agaatgtatt gggaagtcca ctggacatga ggttcac 178

<210> 409  
 <211> 126  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 409  
  
 gcaatggcgg cggagagctt cctgttcacc tcggagtccg tgaacgaggg gcacccagnc 60  
 aagctgttcg ancaggtgtc tgangcggtc tggangcctt cctgnntcag gancccgaca 120  
 ntaaag 126

<210> 410  
 <211> 132  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 410  
  
 gacctcaaga ngggcggcaa caggttcac 60



gacgacgccg acttcacctg ggaggtggtc aagcccctaa agaaggcatc cgcttaagaa 120

tgtattggga ag 132

<210> 411

<211> 83

<212> nucleic acid

<213> Zea mays

<400> 411

gtcggangcg gtgctggang cctgcctggn gcagganncc ganagcaagg tggcctgcga 60

ganctgcacc aagangaaca tgg 83

<210> 412

<211> 133

<212> nucleic acid

<213> Zea mays

<400> 412

gcctcgaccg gatctcgtcg gactcggatc cgcccgacca ccccgcgccg ccgcagatca 60

aagaagatgg cagctgtcga cacattcctc ttcacctcgg agtctgtgaa ngaggganac 120

cctgacaagc tct 133

<210> 413

<211> 290

<212> nucleic acid

<213> Zea mays

<400> 413

tccgatctga gacgagacga gacggnnnnc cctcccctca accggaactt gttttacccc 60

atctcatccc agtgantcnt accacncanc cgcgcgngc ntccgcnnga tctngtcgga 120

ctcggatccg cccgaccacg accaccccgc gtcgccgccg cgcagagcag cagatcagag 180

aagatggccg gactcgacac cttcctcttc acctcggagt ccgtgaacga gggacaccct 240

gacaagtctg cgaccaggtc tcagatgtgt ttggacgttg nttgctgagg 290

<210> 414

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 414

aacaggttca tcaagaccgc cgcatacggc cactttggcc gtgacgacgc cgacttcacc 60  
tgggaggtgg tcaagcccct aaagaaggca tccgcttaag aatgtattgg gaagttcact 120  
ggacatgagg ttcattctcg tctggctctg ctgatacctg caaggatnnn nnnnnnnnnn 180  
nnnnnnnnnn gatgtgtggt tgatcagtga ctggctgctc tgctccatag aagatgaatg 240  
aagagagaga tgggtgaagaa ggctttggca aatggcaatt gccgcagcaa gccatgtcgg 300  
cgccactgac 310

<210> 415

<211> 85

<212> nucleic acid

<213> Zea mays

<400> 415

ctcaggggtgt gcatggccac ttcaccaagc gccccgagga gattggagct ggtgaccagg 60  
gacacatggt cgggtatgcg accga 85

<210> 416

<211> 166

<212> nucleic acid

<213> Zea mays

<400> 416

gagcagcagc gcaaggngan ccgccagctt gccccaggtt ggtagancga gcnagaagaa 60  
ggcaatnncg ggggagagtt cctgttcacn tcggagtccg tgaacgangg gcacccagac 120  
aagctgtgcg accaggtntc ggacgcgggtg ctggacncct gcntgg 166

<210> 417

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 417

aagacggcgg cctatggcca ctttgaaggt gacgaccctg acttcacctg ggaggtggtg 60  
aagccactgc aagtcggaga aaccttctgc ctaaggcggc cttttttttc agtaagaagc 120  
ttttggtggt ctgctgtgct taatcatgct tttatanggc ttctacatgt tgtggttctt 180

tcttgatctg caccgcgctt atcgtttctg ttgtactgcc ctaataagtg gtgcttatga 240  
ggactgtttc tggttttgct gcttatg 267

<210> 418  
<211> 273  
<212> nucleic acid  
<213> Zea mays  
  
<400> 418

acgaccccgga cttcacctgg gaggtggtga agcccctcaa ggcggagaag ccgtcttctg 60  
catgaggcgc ctctctgtgt ttggaagaag cttttggtct ggtctggtct ggtctggtgt 120  
gootgcgcgc tatcatgctt ttttatggct cctacttgtg attcttgatc tgcccccttgc 180  
ttatcatttg tactgtactg tcaactgtcct aataagtggc acgtgtgcgg ggtcgtattg 240  
tgtctgctta ttcacctaga ggattatttc tgg 273

<210> 419  
<211> 57  
<212> nucleic acid  
<213> Zea mays  
  
<400> 419

atcgtctgtg acctgaagga gcatgtcatc aagcctgtca tccctgagca gtacctt 57

<210> 420  
<211> 235  
<212> nucleic acid  
<213> Zea mays  
  
<400> 420

gtcggatctg agacgagacg nngnnncct cccctcaacc ggaacttggt ttaccccatc 60  
tcatcccaact gactcngncc acccaccann ncantgcctc cgccggatct cgtcggactc 120  
ggatccgccc gaccacgacc accccgcgcc gccgccgcgc acagcagcag atcagagaag 180  
atggccggac tcgacacctt cctcttcacc tcggagtccg tgaacgaggg acacc 235

<210> 421  
<211> 297  
<212> nucleic acid  
<213> Zea mays

<400> 421

gccaaagggat gatcaacaat ccaacntcga nctccaagaa ngggcggnaa caggttcac 60  
aagaccgccc catacggcca ctttggccgt gaacgacgcc gacttcacct gggaggtggt 120  
caagccccta aagaaggcat ccgttaagaa tgtattggga agttcactgg acatgaggtt 180  
catcttcgtc tggctctgct gatacctgca aggatnnnnn nnnnnnnnnn nnnnnnnnga 240  
ttgtgtttga tcagtgactg gctgctctgc tccatagaag atgaatgaag agagaga 297

<210> 422

<211> 88

<212> nucleic acid

<213> Zea mays

<400> 422

caencacgag accgtcacca acgacgagat cgccgccgac ctcaaggagc acgtcatcaa 60  
gcccgtgatc cctgagaagt acctgcga 88

<210> 423

<211> 285

<212> nucleic acid

<213> Zea mays

<400> 423

ccgggtcgga tctgagacga gacgagttac catctcatcc caactccgga acgaacaagt 60  
taccatctca tcccaactcc gcctcgaccg gatctcgtcg gactcggatc cgcccgacca 120  
ccccgcgcng ccgcgatca aagaagatgg cntcgtcgac acattcctct tcacctcgga 180  
gtctgtgaac gagggacacc ctgacaagtc tgtgaccagg tctcagatgc cgttcttgag 240  
cttgcnttgc tgaggaccct gacagcaagg ttgttgtgag actgc 285

<210> 424

<211> 136

<212> nucleic acid

<213> Zea mays

<400> 424

accacgacca ccccgcgctc ccgccgcgca naggcagcaga tcagagnaga tagccggatc 60  
tcgacacnt cctcttcacc tcggagtccg tgaacgaggg acaccctgac aagctctgcg 120

accaggtctc agatgc

136

<210> 425  
<211> 217  
<212> nucleic acid  
<213> Zea mays

<400> 425

cgagacgagt nncctcccc cacctcgctt caccacaaccg gaacgaacaa gttacaatac 60  
tcatacccaac cccgccttcg accggatctc gtcggactcg gatccgccc accaccccgc 120  
gccgccgcag atcaaagaag atggcagctg tcgacacatt cctcttcacc tcggagtctg 180  
tgaacgaggg acacctgac aagctctgtg accaggt 217

<210> 426  
<211> 231  
<212> nucleic acid  
<213> Zea mays

<400> 426

cggatctgag acgagacgag ttaccatctc atcccaactc cggaacgaac aagttaccat 60  
ctcatoccaa ctccgcctcg accggatctc gtcggactcg gatccgccc accaccccgc 120  
gccgccgcag atcaaagaag atggcagctg tcgacacatt cctcttcacc tcggagtctg 180  
tgaacgaggg acacctgaca agctctgtga ccagggtcaa tgccgttctt g 231

<210> 427  
<211> 85  
<212> nucleic acid  
<213> Zea mays

<400> 427

agtacctcga ngagaagacc atcttcacc tcaaccgctc cgggcgcttc gtcacgggn 60  
ggntcgangg tgacgtnggc ctcat 85

<210> 428  
<211> 142  
<212> nucleic acid  
<213> Zea mays

<400> 428

caaccccgcc tcgaccggat ctgctcggac tcggatccgc ccgaccaccc cgcgccgccg 60  
cagatcaaag aagatggcag ctgtcgacac attcctcttc acctcggagt ctgtgaacga 120  
gggacaccct gacaagctct gt 142

<210> 429  
<211> 151  
<212> nucleic acid  
<213> Zea mays

<400> 429

cgttcgcctc ttctcctccc tcttgccggg tccttaataa agagcagcag cgcaagaggt 60  
tggtagagcg agcgagaaga aggcaatggc ggcgagagct tcctgttcac ctcgaggtcg 120  
gtgaacgagg ggcacccaga caagctgtgc g 151

Sequence

<210> 430  
<211> 257  
<212> nucleic acid  
<213> Zea mays

<400> 430

agtgtcctc ctctatgaa gaaggactct ccacgttctc nccgtcaaag cagtggaaaa 60  
cagatttggg cgaggtagca acagcgtctt cagcagcgca gaagctctgg tatacgagga 120  
gcccgcctcc aggcagctcc tgctccacca tggtgttgca gtcatagacc aaatctgttt 180  
tccactgctt tgacggcgag aantgganna gtgctcctcc tcctatgaag aaggactaca 240  
aactagctaa tcttctc 257

<210> 431  
<211> 220  
<212> nucleic acid  
<213> Zea mays

<400> 431

aaagagatga cgaagctctc ggggattcat gagatcattc ccgagatgga gatctgtgac 60  
tttgagtttg acccctgtgg gtactcgatg aatggcgtct tcgggcctgc agcctocacc 120  
atccacgtga cacccgagga aggtttcagc tacgcaagct acgaagctat gaacttcgac 180  
cccagctcac tgggtctacag cgatgtgatc aggagggtcc 220

<210> 432  
 <211> 240  
 <212> nucleic acid  
 <213> Zea mays

<400> 432

gggagaaatt cgtgagatct tggnnctgnt cagggcgtgc gagcncctgga atcatggggtt 60  
 tcacacatag cttcgnctnt tngaatttna tgtactaatg gagtcnaagg gtggcaaaaa 120  
 gtcnnncngt agtcgttcta tgggtgatga agcgccctt ggctacagca ttgaggncgt 180  
 tcgacctgcc ggagcgtgaa gaantccagc tgcggcttac tcgaactgcg ngaatnagcg 240

<210> 433  
 <211> 263  
 <212> nucleic acid  
 <213> Zea mays

<400> 433

cttgtgtcgg tagttttccg ttggctctgg ctgccttctt ctgcttctga gattccaact 60  
 tgtcttgccc atctctcctt ttcctctctc atcctctctt tcttgcaaca cgtcatccag 120  
 tggcgatgtc ttcagcagat tcttgtgtct cttctcctgc ctccctatt ggctttgagg 180  
 gctatgagaa gcgcctcgag atcacgttct ctgacgcgcc tgtctttgag gacccttggtg 240  
 gtctgtgcnt gcgcgcctc tcc 263

<210> 434  
 <211> 290  
 <212> nucleic acid  
 <213> Zea mays

<400> 434

taatgtgat ggcaacacaa cattagtctt gaagaagaat gaagctttct tcaagactaa 60  
 tgctgatggc aacacaacat anttccaagg aaatgacgaa gctctctggt atctctgaaa 120  
 ttatccctga gaaggagatc tgtgattttg acttcgaacc ctgcggctac tccatgaatg 180  
 caatccatgg ctctgcgttc tccacgatcc atgtgacgcc tgaggatggg ttcagctacg 240  
 ccagttatga ggttatgggc ttggatgcca ctgccctgtc ttacggtgac 290

<210> 435

<211> 258  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 435  
  
 tggagcagga gctgcctgga ggcgggctcc tcgtgtacca tngcttctgt gntgctgaag 60  
 acgtgtttgc tacctcgccc aaatctgttt tccacngcnt tganggcgag aacgtggaga 120  
 gtgtcctctn tcgnntgagg agganancaa gnggnnaang ttctntgttg ggagnatnan 180  
 acggngtgcc atggaggaga nggcgggatt ccttgatgag taatangggg ctctgggntc 240  
 gattagcttc tgattgtt 258

<210> 436  
 <211> 263  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 436  
  
 cttgctctaa aacaatcgca gtcttgcaag ttgttgctgc tgctcctacc cctgcctctg 60  
 caattgggtt tgagggatat gagaagcgcc tcgagatcag cttctatgag gcacctgtct 120  
 tcgtgaccc caacggaagg ggattgtgtg cactcttgcg tccctagatt tactctatctt 180  
 ctgaccatgc acggtgcacc gttgtctctg agctatcaaa cgaggacttt gactctatgt 240  
 cttatctgag tcaagcctgt ttg 263

<210> 437  
 <211> 266  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 437  
  
 cccaganatg gagatctgtg acttcgactt cnagccctgt ggctactcca tgacatgctg 60  
 ttcattggcnc tgcnattgtc gaccattcat gnganccnc aggacgggca tcagctaattg 120  
 aaagcaacan ggncatgggc nttaanccgg ggctcctttc tcatatggtn anctggntaa 180  
 ganggtgctg aagnnntttn gncnactga ntncnctgtt gncgtaacc atcttccggt 240  
 gatcgggcaa tgcgaagacc tggggg 266

<210> 438



<211> 281  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 438  
  
 gggctgcccg gtgtaggcaa gtcaaccatc atgagcgagc tttattaata tgatgcaagc 60  
 tacctggttt ggtggtaatg cttatgtgat tggtgattct gcaaagcata agcagaagtg 120  
 gcacgtctac tatgccacca ctgagcacc ctaggagct tgttgttact cttgagatgt 180  
 gcatgactag gctggacaag aagagagctt atgtcttctt caagacctct attgatgggt 240  
 acacatcttg tgctaaggat atgaccaagc cttcagtgat t 281

<210> 439  
 <211> 334  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 439  
  
 gcagaagtgg cacgtctact atgccaccac tgagcacc c taaggagctt gttgttactc 60  
 ttgagatgtg catgactagg ctggacaaga agagagctat gtcttcttca agacctctat 120  
 tgatggttac acatcttgtg ctaaggatat gaccaagcct tcagtggcat cgtcaaggag 180  
 tgcgctgtga ccagtgggtg tagcagcaag gccatggcac tcagggtcca tgggtcgtgg 240  
 gtgtggcttg ttcagcgctt actattcgcg ggagtcggcg aggtgcctgc tgaacttggt 300  
 ggaagtggag tccattatga tctgacacga cctc 334

<210> 440  
 <211> 349  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 440  
  
 catctatcct ctgaagattg tcatcaagac ctgtggcact accaagctcc tgctcacaat 60  
 tocaaggatc ctagagcttg tgaagagctg tctatgctct tgtgctgtga antatcccgc 120  
 gggacgttca tctttcttg cgnacagcag cccccaccg gagttctccg aggagttgtg 180  
 tattaaccgt actttggggg ctgaagtctg gtggcatgct tatgtgattg gagatgcagc 240  
 aagaccagga cagaagtggc acatctatac gccactgagt acccagagca accatgtcac 300

cttgagatgt gcatgatggt tggacagaag aagcttcac tttttcaag

349

<210> 441  
<211> 260  
<212> nucleic acid  
<213> Zea mays

<400> 441

tttctcttat ggngacctgg ntaagagggt gctgaggcgc ttnggtccaa ctgagcnctc 60

tgttgccgtg accatottcg gtgatcgga caatgcgaag acctggggga cgaaactgna 120

tgctgaggcc tatgcttgca gcaacatggc tgagcaggag cttgccgatt ggtggcttgc 180

tcatttatca gagcttcaact gttacggcgc aancgacctn tgnttcccag ngcaaacntc 240

gnaaattccg ggnaaantaa 260

<210> 442  
<211> 447  
<212> nucleic acid  
<213> Zea mays

<400> 442

cgccctgca caggcggtt ttcagactgc gatccattcc gagcttcgag gaatcgatgg 60

tcnagcttcg gtgactaaga gatcaaactc ttcaagttct acgaggctga agatcctgag 120

catctgtttg gtgaagattc ttatgccatg gaaattcatc gattgatacc gtccctttca 180

tagtcgatca tggttcaagg agattgcatt tgtggatgct ctaatggagt cgaaaggtgg 240

taagaaagtc nnnnnnnnnn nnnnnnnnnt tcatgtacga agctcccctt ggctacaaga 300

tcgaggacgt tcgcccagcc ggaggaatca agaagttcca gactgctgct tattccaact 360

gcgtccgcca gccatcctga tatcgctca catgcaattc gcggtagagt aggattttaa 420

ttcagttttc ctcttngtc ngnaggt 447

<210> 443  
<211> 192  
<212> nucleic acid  
<213> Zea mays

<400> 443

gggctgagac ttcgannacg agccctnttg ctattncatg aatgctgnaa atnanccggg 60

gttggggana attcatntga gncnagagga cngattnagc tatgcaagct annaggtcat 120  
 gggctngaac cccgactctt tntgttatgg tgacctgcct aanagggagc tgangngctn 180  
 nggtccaaat ga 192

<210> 444  
 <211> 376  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 444

caaagctnnt gctcgctatt ccgaggntcc ttgnactngn tgaanagctc cngttgcnac 60  
 tcgctgcagt taaanactnn cgtggaacgt ncatattccn tgaagcacan centtcnnac 120  
 acaagaactt ngntgacgag gttgccttcc tgaatngctt attcngtggn ctnaagtnen 180  
 gnngcaatgc ttatgtgant ggtgactctg ccaancccg gncagaagtgn cacgtctact 240  
 acacnnctga gcacctgan gagcctgttn gnactctgga gatgtgcatg actgggctgg 300  
 acaagancaa agcttcatgn cntcttcaag accactgnnt gatggttact cgctnatncc 360  
 ccaaggatat caccaa 376

<210> 445  
 <211> 502  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 445

gtgctagtgg cnangnnntn nngtatnaan gacacactcc ggtncggaat tcccgggtcg 60  
 acccacgcgt ccgcgagcgc gtgggggtgc caaagaattn cctnccctac cgtcgctcgc 120  
 gctcgctctgg nngggaagtt ccnggaccn ngatttngcc caattctnag gnaaaaaann 180  
 ggtttacntn ncttaaccn gggnnctctn cctntngtng ttgttaaagg ganaaaattn 240  
 tnagatctgt tccgatcaa gcgtgcganc tcgggaatca agggttttna cacatatctt 300  
 ttccatttga aaattgatnt actaaatggn gtctanaagg tggcaaaaag tctnacaata 360  
 gtcgttctat natgtatgaa gctccccttn gctncagcat tnaggacgtt cgacctgcc 420  
 gangcctnaa gaaatttcca gtctnctgct tactncaact tcncaaaaaa accattctga 480  
 tatecttttg cttnctcaat nt 502

<210> 446  
 <211> 160  
 <212> nucleic acid  
 <213> Zea mays

<400> 446

agtnngctgca cttaaccggt nctttggcgg cctgaaatct ggtggnangg gcgggtgtga 60  
 tnggagatcc agcaagacct ggncagaagt ggcacttntt nnaacgnan tnantnccna 120  
 ttancaacca atggttaacc tanaaatgtg catnactgna 160

<210> 447  
 <211> 487  
 <212> nucleic acid  
 <213> Zea mays

<400> 447

attacgtccn ctagcccttc gccnncgctc ttttccntna gaggcgccgg caccgangat 60  
 ccgggncggg ntcnatntnn agaannngna gtacttgtct cagcccgggg nctgctgcgt 120  
 cnggtggtgg tnaaggggag aanntcgtna gatctgntcc ggatcaagcg nncganctcg 180  
 ggaatcaggg gtttcacaca tanctgngnc gatcnogaat tctgatgtac taatggagtc 240  
 taagggnggc aaaaagtcta gccttacnta cnctatnatg catagaagct ccccttggtc 300  
 acagcattna cgaccgttcg anctgtccng aggcgtgaaa aagntccnnt acngctgctt 360  
 actccaactt gcgcgaaaca anccntcctg aanantcccc ttttggtctc ctcattctaa 420  
 gcactttaag gaattttaat ntctggacac ttntggantn ttnaccaaan ctentctggg 480  
 cctaggg 487

<210> 448  
 <211> 438  
 <212> nucleic acid  
 <213> Zea mays

<400> 448

gagganntcg ttgagatcta agatagagaa tcgctggggc angngnctgt ggtctgctgc 60  
 gtaggggtggt gntgaanggg agaagtttaa gatctgntcc acagatcacg cgtgcgcgct 120  
 cgcgaancgg gggttcacac catagcttng tcggtttgaa tttgangtac taatggagtc 180

taaggggtggc aagaantcta gcagnagtcg ttccatgatg natgaagctc cccttggtcta 240  
cagcattgaa gacgttcgac ctgccggaag cgccaaagaa gttncagnct tgntgcttaa 300  
ctccaactgg cncaaaagaa nccatnctgn aatccccctt ggnttccccg ntctangagn 360  
ttaagaattc ttttcttgca ctttnaatct taacnaatcc ccctgggctg angnttttct 420  
gacaaaangaa ccaattnc 438

<210> 449  
<211> 429  
<212> nucleic acid  
<213> Zea mays  
<400> 449

ctcgattaca aactgtgtac tatgttaatg taatgacata gtttggtga tacagatcgt 60  
agtgatgttg tttgaagctg tttgtatnaa ctgtntccta ctatcttgta tgaattatga 120  
tgtctgatgt agtntgtatg aactatgtca tattatgttg aatgaagcta ttgcatgccc 180  
ttaaaaaana aaaaccnaag annanaatga tctactccaa ntaacgagnt tatgggcttg 240  
gatgccactg ntntgnntta tgggtgacct gtcaagaggg tgctnctggn ctttngccnn 300  
tnagagtttt acnttgacct gaccatcttc nncggncgtg gnatgccggg acatnnggna 360  
aggnacttgg tgnanaagtc tttnactggn acanctgtt cgaacangac ttntnttnaa 420  
gcggnctcc 429

<210> 450  
<211> 376  
<212> nucleic acid  
<213> Zea mays  
<400> 450

ccaatatctt tactnccgcc cctgncgta attccctnct ggcncacgc atcngctga 60  
agacnctn tnctacctn ccnantntg tttccactn ctttnaccac gagggngggg 120  
nagattgctc ctntctctat caataangac tacaatctgg ctaatcttct ctgctnaaca 180  
ngaggaatcc ccatgccatn natnaccaag ncnntagtgc ttgatnanta atacntctt 240  
ctaagntcca tntncttctg aattgnntat antatctct cacantttca tantntcan 300  
tnagttattc tatcaagcat ccaatccatt ctattgtata ttaaaatttn tctctctnta 360

tncatgtcaa cttcct

376

<210> 451  
<211> 305  
<212> nucleic acid  
<213> Zea mays

<400> 451

ggaagttgga tgctatcgag gagaaggatg gagtgcttga tgagtaagac gggtttctgg 60  
tgtcgatttg cttctgagtt tttttctnnt ttatatcggt gcgatctcgt ggttgctcgt 120  
tggttagcag ccaagccagg ctattagtag gaagaatgtc gtctataaac atgtgttcct 180  
acgttgccac atgtcgagtc aatctgtata cagctagctc taggtgggtca gctgcgtcta 240  
ccacaatgag catacatgta tggataaatc ttctgtgaaa gcattcgatg aataagggtt 300  
gtttt 305

<210> 452  
<211> 276  
<212> nucleic acid  
<213> Zea mays

<400> 452

atgcaagcac cttttgcgaa ggactgcaag ctggctaata ttgtctgctt ggaggaagtt 60  
ggatgctatc gaggagaagg atggagtgtc tgatgagtaa gacgggtttc tggctcgatt 120  
tgcttctgag tttttttctt tttatatcgt tgcgatctcg tggatgtcgt ttggttagca 180  
gccaagccag gctattagta cgaagaatgt cgctataaac atgtgttcoo acgtgccaca 240  
tgctgagtca atctgtatac agctantcta ggtggc 276

<210> 453  
<211> 501  
<212> nucleic acid  
<213> Zea mays

<400> 453

gnaagngttt tgaatggggg ntgggngna annnnnnnct ngtaaccggtc gngngnctgn 60  
cccacgnta cgcgccnttg ntagctcggc ccaaatacgt tcttngntgc tttgatggcg 120  
agaatgcagc accttttgcg aaggactgca agctggctaa tcttgtctgc ttggaggaag 180

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 tcgtntt 247

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gtcgtctata aacatgtgtn cctacgttgc cacatgctga gtcaatctgt atacagctag 60  
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338

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<400> 461

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aaaggtggca agaagtcnnn nnnnnnnnnn nnnnnnttca tgtacgaagc tccccttggc 240

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<210> 463



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<400> 488

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<210> 489

<211> 304

<212> nucleic acid

<213> Zea mays

<400> 489

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ctgaaacgat ggcagtcctg caagttgctg ctgctgcccc tccccctgtc tctgtgattg 180  
ggttcgaggg atttgagaag cgccttgaga tcagcttctc tgaggcacct gtcttggtg 240  
accccagcgg aaggggactg cgtgcgctct cgcgtgccca gatcgactct gttcttgacc 300  
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<210> 490

<211> 229

<212> nucleic acid

<213> Zea mays

<400> 490

gaagacggga agtgggtcttt gatattcttt ggtcactgga acgatggcag tcctgcaagt 60  
tgctgctgct gcccctcccc ctgtctctgt gattgggttc gagggatttg agaagcgct 120  
tgagatcagc ttctctgagg cacctgtctt ggctgacccc agcgggaagg gactgcgtgc 180  
gctctcgcgt gccagatcg actctgttct tnaattgctc ggtgcacca 229

<210> 491

<211> 290

<212> nucleic acid

<213> Zea mays

<400> 491

gctgaggtgc tttggtccaa ctgagttctc tgttgccgtg accatcttcg gtgatcggga 60  
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ttgagcaggt gctgccgttt ggtggcttgc tcatctatca gagcttcact gttacggccg 180  
aaacgaccca tgggtcgccg aggtcagtec tgcattgactt cgctgggtgac attgtcaaga 240  
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<210> 492

<211> 289

<212> nucleic acid

<213> Zea mays

<400> 492

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gacctgtgga actacaaagc tatctgctcg ctattccgag gatccttgaa cttggctgaa 120  
gagctcctgt tgccactcgc tgcagttaaa tactcccgtg gaacgttcat attccctgaa 180  
gcacaacctt cccaacaca agaacttcgc tgacgagggt tgcctnctg aatcgctct 240  
cgggtgggctc aangtcgggtg ggaaagctan tgtgattgtn annnngcga 289

<210> 493

<211> 318

<212> nucleic acid

<213> Zea mays

<400> 493

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caatgcgaag acctggggga cgaaactgga tgctgaggnc tatgcttgca gcaanctggt 120  
tgagcaggtg ctgccgtttg gtggcttgct catctatcag agcttcactg ttacggccga 180  
aacgacccat gggtcgccga ggtcagtcct gcatgacttc gctgggtgaca ttgtcaagaa 240  
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tgtaaagaag atgnaatg 318

<210> 494

<211> 310

<212> nucleic acid

<213> Zea mays  
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 tacctccggg tagctgcttt ccagtgacca aaagatatca aagaccactt cccgtcttcc 120  
 tcccgtggct gtctgtcatc tggctctgaa acgatggcag tcctgcaa at tgctgtgct 180  
 gccctcccc ctgtctctgt gattgggttc gagggatttg agaagcgcct tgagatcagc 240  
 ttctctgagg cactgtcttg ggctgacccc agcgaaggg actgcgtgcg ctctcgcgtg 300  
 cccagatoga 310

<210> 495  
 <211> 137  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 495  
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 tttgggtcaa ctgagttctc tgtcaccgtg accatcttcg gtgatcggga caatgcgaag 120  
 acctggggga cgaaact 137

<210> 496  
 <211> 111  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 496  
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 acaatgcgaa gacctggggg acnnaactgg atgctgaggc ctatncttag c 111

<210> 497  
 <211> 302  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 497  
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atggagtcca aggggtggcaa gaagaagtct agcagtagtc gttcctccct gatgtacgaa 180  
gctcccccttg gctacagcat cgaggatatc cgccctgcag gcggcatcaa gaagttctcc 240  
gctgcttact cgaactgcg c gaggaagcca tctgatatac tctgtcgtca tccccatcct 300  
ag 302

<210> 498  
<211> 501  
<212> nucleic acid  
<213> Zea mays  
<400> 498

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tagcaccgaa gagaaatata tgccgcgcgc tctcctcat tcggtgaaag gtgggcggtt 120  
agggttggga agatctcttg actcagaca cgagcagcag aattcgtcga gggccgctgg 180  
ttgctgcgag aggtccgggc ctctgtggga tcttgaaaat tgaatgttct aatggagtcc 240  
aagggtggca agaagaagtc tagcagtagt cgttcctccc tgatgtacga agctccccctt 300  
ggctacagca tcgaggatat ccgccctgca ggccgcatca agaagttctc cgctgcttac 360  
tcnaactgcn cgaggaagcc atcctgatata ctctgtcgtc atccccatcc taatagccgt 420  
naaaaccanc tggaatttca atttgatctn cccaaaatcc ctcgggttnc cggctttcca 480  
gtgaccanag ngatatcaaa g 501

<210> 499  
<211> 284  
<212> nucleic acid  
<213> Zea mays  
<400> 499

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gacaaaaaga tatcagagac cacctcccgt ggctgtctgc catctggctc tgaaacgatg 180  
gcagtccctac aagctgctgc cctccccct gtctctgtga ttggattcga gggatttgag 240  
aagcgccttg agatcagctt ctctgaggca cctgtcttgg ctga 284

<210> 500

<211> 166  
 <212> nucleic acid  
 <213> Zea mays

<400> 500

caagaagaag tctagcagta gtcgttcttc cctgatgtac gaagctcccc ttggtacag 60  
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 cgcgaggaag ccatacctgat atctctgtcg tcatacccat cctagt 166

<210> 501  
 <211> 440  
 <212> nucleic acid  
 <213> Zea mays

<400> 501

gagatctctt gctcacgac acgagcagca gaatttgtcg agggccgggtg gttgctgcga 60  
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 aagaagaagt ctagcagtag tegtctcttc ctgatgtacg aagctccctt tggctacagc 180  
 atcgaggata tccgctgca ggcggcatca agaagttctc cgctgcttac tcgaactgcg 240  
 cgaggaagcc atcctgatat ctctgttgtc atcccatcc taatagccgt agaaaccacc 300  
 tgcattttca ttttgatctc ctaatccctt cggctagctg cttttcaagt gacaaaaaga 360  
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 cgatggcagt cctgcaaggt 440

<210> 502  
 <211> 300  
 <212> nucleic acid  
 <213> Zea mays

<400> 502

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 gaggaagcca tctgatata tctgtcgtca tccccatcct agtagcgtag aaacctctg 180  
 cattttcatt ttgatctccc taatctctcc ggctagctgc tttccagtga ccaaaagata 240  
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<210> 503  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 gctagctgct ttccagtgac caaaagatat cagagaccac ctcccgtggc tgtctgccat 120  
 ctggctctga aacgatggca gtcttacaag ctgctgcccc tccccctgtc tctgtgattg 180  
 gattcgaggg atttgagaag cgccttgaga tcagcttctc tgaggcacct gtcttggtg 240  
 accccagcgg aaggggactg cgtgcgctct cgcgtgccca gat 283

<210> 504  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 ggcggttagg gttgggagat ctcttgactc acgacacgag cagcagaatt cgtcgagggc 120  
 cgctgggttc tgcgagangt cggggccttc gtgggatctt gaaaattgaa tgttctaattg 180  
 gagtccaagg gtggcaagaa gaagtctagc agtagtcgtt cctccctgat gtacgaagct 240  
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<210> 505  
 <211> 313  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 505  
  
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 gcagcagaat tcgtcgaggg ccgctgggtt ctgagagang tccgggcctt cgtgggatct 180  
 tgaaaattga atgttctaatt ggagtccaag ggtggcaaga agaagtctag cagtagtcgt 240  
 tctccctga tgtacgaagc tccccttggc tacagcatcg aggatatccg ccctgcaggc 300

ggcatcaaga att

313

<210> 506  
<211> 283  
<212> nucleic acid  
<213> Zea mays

<400> 506

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ggcgggttagg gntgggagat ctcttgactc acgacacgag cagcagaatt cgtcgagggc 120  
cgctgggttg tgcgagangt ccgggccttc gtgggatctt gaaaattgaa tgttctaata 180  
gagtccaagg gtggcaagaa gaagtctagc agtagtcggt cctccctgat gtacgaagct 240  
ccccttggt acagcatcga ggatatccgc cctgcaggcg gca 283

<210> 507  
<211> 304  
<212> nucleic acid  
<213> Zea mays

<400> 507

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tcctcattcg gtgaaaggng ggcgggttagg gttgggagat ctcttgactc acgacacgag 120  
cagcagaatt cgtcgagggc cgctgggttg tgcgagaggt ccgggccttc gtgggatctt 180  
gaaaattgaa tgttctaata gagtccaagg gtggcaagaa gaagtctagc agtagtcggt 240  
cctccctgat gtacgaagct ccccttggt acagcatcga ggatatccgc cctgcaggcg 300  
gcat 304

<210> 508  
<211> 439  
<212> nucleic acid  
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<400> 508

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ctcctcctca ttccgtgaaa ggtgcgcagt taggggtggg agatctcttg cctcacgaca 120  
cgagcagcag aatttgctga gggccgctgg ttgctgcgag aggtccgggc cttcgtggga 180

tcttgaaaat tgaatgttct aatggagtcc aaggggtggca agaagaagtc tagcagtagt 240  
 cgttcctccc tgatgtacga agctccccctt ggctacagca tcgaggatat ccgcctgcag 300  
 gcggcatcaa gaagttctcc gctgcttact cgaactgcgc gaggaaacca tinctgatatc 360  
 tctggtgnca ttcccattct aatagccgta gaaaccacct gnattttcaa tttgaacttc 420  
 ccttttaaaa atccttccg 439

<210> 509  
 <211> 453  
 <212> nucleic acid  
 <213> Zea mays

<400> 509

cgcggtggttc gttaggaaag gaaatgggat tccccagca tctgatctct tcggtgctag 60  
 agaaatatan gccggcgncg cctnctactc attcggtgaa aggtgcgcag ttaggggttg 120  
 gagatctctt gcctcacgac acgagcagca gaatttgtcg agggccgntg gttgctgcga 180  
 gaggtccngg ccttcgtggg atcttgaaaa ttgaatgttc taatggagtc caaggggttgc 240  
 aangaaagaa atctagcagt agtcgttctt tcttgatgta cgaaacttcc nttggetnca 300  
 acatcgagga tattcgccct tgaagcngca ttaaanantt nttcnntngt tactnnaant 360  
 ttncangaa accattctta aatnctttgt ngnattccca tcctaatacg tttnaaacan 420  
 ctgaatttca attgacttct naaaanctag ggt 453

<210> 510  
 <211> 261  
 <212> nucleic acid  
 <213> Zea mays

<400> 510

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 gggcggttag ggttgggaga tctcttgact cacgacacga gcagcagaat tcgtcgaggg 120  
 ccgctgggtt ctgcgagang tccgggcctt cgtgggatct tgaaaattga atgttctaata 180  
 ggagtcnag ggtggcaaga agaagtctag cagtagtcgt tcctcctgat gtacgaagct 240  
 ccccttggtt acagcatcga g 261

<210> 511

<211> 190  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 511  
  
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 atcctgatat ctctgtcgtc atccccatcc tagtagcgta gaaacctcct gcattttcat 120  
 tttgatcncc ctaatctctc cggctagctg ctttcagtg accaaaagat atcagagacc 180  
 acctcccgtg 190

<210> 512  
 <211> 307  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 512  
  
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 gaaatatagg ccgccgccgc ctctccttg ttcggtgaaa ggtgggcagt tagggttggg 120  
 agatctcttg actcacgaca cgagcagcag aattcgctga gggccgctgg ttgctgcgag 180  
 aggtccgggc ctctgtggga tcttgaaaat gaatgttcta atggagtcca agggtgggca 240  
 gaagaagtct agcagtagtc gttcctccct gatgtacgaa gctccccttg gctacagcat 300  
 cgaggat 307

<210> 513  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 513  
  
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 ctctccttg ttcggtgaaa ggtgggcagt tagggttggg agatctcttg actcacgaca 120  
 cgagcagcag aattcgctga gggccgctgg ttgctgcgag aggtccgggc ctctgtggga 180  
 tcttgaaaat tgaatgttct aatggagtcc aagggtggca agaagaagtc tagcagtagt 240  
 cgttctctcc tgatgtacga agtcccctt ggctacagca tcgaggatat ccgccctgca 300  
 gcggcat 307

<210> 514  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 514  
  
 agaaatcgat tcccccgcat cggatctctt cggtgctaga naaatatagg ccgccgcctc 60  
 ctcttgcatt ncggtgaaag gtgggcgggt aggggttgga gatctcttga ctacacgacac 120  
 gagcagcaga attcgtcgag ggccgntggg tgctgcgaga ggtccggggc ttcgtgggat 180  
 cttgaaaatt gaatgttcta atggagtcca aggggtggca gaagaagtct agcagtagtc 240  
 gttcctccct gatgtaogaa gtccctggc taca 274

<210> 515  
 <211> 290  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 515  
  
 aaatcgattc ccccgcatcc gatctcttcg gtgctagaga aatataggcc gccgcctcct 60  
 cctcattcgg tgaaagggtg gcgggttaggg ttgggagatc tcttgactca cgacacgagc 120  
 agcagaattc gtcgaggggc gctgggttgc gcgagaggtc cgggccttcg tgggatcttg 180  
 aaaattgaat gttctaattg agtccaaggg tggcaagaag aagtctagca gtagtcgttc 240  
 ctccctgatg tacgaagctc cccttggtc cagcatcgag gatatccgcc 290

<210> 516  
 <211> 291  
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 <213> Zea mays  
  
 <400> 516  
  
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 gcgggttaggg ttgggagatc tcttgactca cgacacgagc agcagaattc gtcgaggggc 180  
 gctgggttgc gcgagaggtc cgggccttcg tgggatcttg aaaattgaat gttctaattg 240  
 agtccaaggg tggcaagaag aagtctagca gtagtcgttc ctccctgatg t 291

<210> 517  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 aggat 305

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 cagaattcgt cgagggccgc tggttgctgc gagaggtccg ggccttcgtg ggatcttgaa 180  
 aattgaatgt tctaattggag tccaaggggtg gcaagaagaa gtctagcagt agtcgttcct 240  
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 <212> nucleic acid  
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 ttgactcacg acacgagcag cagaattcgt cgagggccgc tggttgctgc gagaggtccg 180  
 ggccttcgtg ggatcttgaa aattgaatgt tctaattggag tccaaggggtg gcaagaagaa 240

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<210> 520

<211> 287

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<213> Zea mays

<400> 520

aaggnaatcg atnccccgc atccgatcnc ttcggtgcta gagaaatata ggccgcccgc 60

tccnccatcat tcggtgaaag gngggcggtt agggttggga gatctcttga ctacacgacac 120

gagcagcaga attcgtcgag ggccgctggt tgctgcgaga ggtccggggc ttcgtgggat 180

cttgaaaatt gaatgttcta atggagtcca aggggtggcaa gaagaagtct agcagtagtc 240

gttccctccct gatgtacgaa gctccccttg gctacagcat cgaggat 287

<210> 521

<211> 298

<212> nucleic acid

<213> Zea mays

<400> 521

gaaatcgatt cccccgcac cgatctcttc ggtgctagag aaatataggc cgccgcctcc 60

tcctcattcg gtgaaagggtg ggccggttagg gttgggagat ctcttgactc acgacacgag 120

cagcagaatt cgtcgagggc cgctggttgc tgcgagaggt ccgggccttc gtgggatctt 180

gaaaattgaa tgttctaatt gagtccaagg gtggcaagaa gaagtctagc agtagtcggt 240

cctccctgat gtacgaagct ccccttggt acagcatcga ggatatccgc cctgcang 298

<210> 522

<211> 313

<212> nucleic acid

<213> Zea mays

<400> 522

cgcgtgggttc gttaggaaag gaaatcgatt cccnccgcat ctgatctctt cgggtgctaga 60

gaaatatagg ccgccgccgc ctctccttg ttcggtgaaa ggtgggcagt tagggttggg 120

agatctcttg actcacgaca cgagcagcag aattcgtcga gggccgctgg ttgctgcgag 180

agggtccgggc cttcgtggga tcttgaaaat tgaatgttct aatggagtcc aaggggtggca 240  
agaagaagtc tagcagtagt cgttcctccc tgatgtacga agctcccctt ggctacagca 300  
tcgaggatat ccg 313

<210> 523  
<211> 265  
<212> nucleic acid  
<213> Zea mays

<400> 523

cgcggtggttc gttaggaaag gaaatcgatt cccccgcac ctgatctctt cggtgctaga 60  
gaaatatang ccgccgcgc ctcctccttg ttccgtgaaa ggtgggcagt tagggttggg 120  
agatctcttg actcacgaca cgagcagcag aattcgtcga gggccgctgg ttgctgcgag 180  
agggtccgggc cttcgtggga tcttgaaaat tgaatgttct aatggagtcc aaggggtggca 240  
agaagaagtc tagcagtagt cgttc 265

<210> 524  
<211> 272  
<212> nucleic acid  
<213> Zea mays

<400> 524

gngtggttcg ttaggaaagg aaatcgatnc nccccgcac tgatcncntc ggtgctagag 60  
aaatataggc cgccgccgcn nccnccctt gttccgtgaa aggnnggcag ttagggttgg 120  
gagatcncctt gactcacgac acgagcagca gaattcncg agggccgctg gttgcngcga 180  
gaggncggg cttcgtggg atcttgaaaa ttgaatgtnc taatggagtn caaggggtggc 240  
aagaagaagt ctagcagtag tcgttcctcc ct 272

<210> 525  
<211> 550  
<212> nucleic acid  
<213> Zea mays

<400> 525

agttataagn ggggntagn aaggaacgct ntnccgtcc ggaattccc ggtcgacca 60  
cgcgtcgcga gcattgagga cgttcgacct gccggaggcg tgaagaagtt ccagtctgct 120



gcttactcca actgcgcgaa gaagccatcc tgatatccct tttggcttcc tcattctagt 180  
 agnttaggat ttcttttctg acactttgat tctgaccaat ctctctggcc tgetgcttcc 240  
 tgataatoga ccagttcccc agtcttgctc cttgcactcc tccctccatc tccagcattg 300  
 tgttctgatt cacctgctcc aatggctgtt ctttctgctg ctgatgcttc ccggtctcag 360  
 ctatcggttt tgagggctat gagaagcgcc ttgagatcac attctctgag gcacctgtct 420  
 ttgtggaccc tcatggggcg tggtttgctg gccctctnca gggcccanat tgactctgtt 480  
 ctggatcttg cacggtgcac aattgtgtnc ganctctnca acaaggattt cgactcatat 540  
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 <211> 537  
 <212> nucleic acid  
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taatgtgntn nnnntggggn tnggggagga nanngtaant gcttctaccg gtccggaatt 60  
 cccgggtcga cccacgcgtc cgcaaagctc tctggcatct ctgaaatcat ccccgagatg 120  
 gagatctgcg attttgactt tgaaccctgt ggctactcca tgaatgcgat ccatggctct 180  
 gcattctcca caatccatgt gacgcccag gacggtttca gctatgccag ttatgagggt 240  
 atgggcttgg atgccactgc tctgtcttat ggtgaccttg tcaagagggt ccttcggtgc 300  
 tttggccctt cgaggttttc cgttgccgtg accatcttcg gcgggctgtg ccatgccggg 360  
 acatggggaa aggcacttgg tgcagaggtc tatgactgca acaacatggt ggagcaggag 420  
 ctgcctggag gcggctcttc gtgtaccaga gcttctgtgc tgctgaagac gctgttgcta 480  
 cctcgcccaa atctgttttc cactgctttg acggcgagaa cgtggagagt gctcttc 537

<210> 527  
 <211> 544  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 527

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 nnacgcgtcc gccacgcgt ccggaagaaa gcttctgtct tttcaagac taatgctgat 120

gggaacacaa catgtgccaa ggaaatgaca aagctctctg gcatctctga aatcatcccc 180  
 gagatggaga tctgcgattt tgactttgaa ccctgtggct actccatgaa tgogatccat 240  
 ggotctgcat tctccacaat ccatgtgacg cccgaggacg gtttcagcta tgccagttat 300  
 gaggttatgg gcttggatgc cactgctctg tcttatgggtg accttgtcaa gagggtcctt 360  
 cgggtgctttg gccctcggga gttttccgtt gccgtgacca tcttcggcgg gcgtggccat 420  
 gccgggacat ggggaaaggc acttgggtgca naggtctatg actgcaacaa catgggtggaa 480  
 cangagctgc ctggaagcgg gctnctcgtg taccaaagct tctgtgctgc tnaaacnctg 540  
 ntgc 544

<210> 528  
 <211> 539  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 528

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 gcttgtttat ctgtcctctg aagattgtca tcaagacctg tggcactacc aagctcctgc 180  
 tcaccattcc aagaatcctt gagcttgctg aagagctgtc tatgccactt gctgctgtga 240  
 agtactcccg tgggacgttc atctttcctg gcgcacagcc agccccccac aggagcttct 300  
 ctgaggaagt tgctgcactt aaccgctact ttggcggcct gaaatctggg ggtaatgctt 360  
 atgtgattgg agatccagca agacctggac agaagtggca cgtctttctac gccactgagt 420  
 acccagagca accaatgggtt aaccttgaaa tgtgcatgac tggctctggac aagaaagaaa 480  
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tgaagtactc ccgtgggacg ttcatctttc ctggcgcaca agccagcccc ccacagggct 180  
tctctgagga agttgctgca cttaacctgt actttggcgg cctgaaatct ggtgggtatgc 240  
ttatgtgatt ggagatccag caagacctgg acagaagtgg cacgtcttct acgccctgag 300  
taccagagac aaccaatggt taaccttgag atgtgcatga ctggtctgga caagagaaaag 360  
cttgtgtctt tttcaagact aatgctgatg ggaacacaac atgtgccaag gaatgacaaa 420  
gtctcttgga atctctgaaa tcctccccga gatggagatc tgcgattttg ac 472

<210> 530  
<211> 421  
<212> nucleic acid  
<213> Zea mays

<400> 530

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cgtgaccatc ttccggcggg gtggccatgc cgggacatgg ggaaaggcac ttggtgcaga 180  
gggttatgac tgcaacaaca tgggtggagca ggagctgcct ggaggcgggc tcctcgtgta 240  
ccagagcttc tgtgtctgtg aagacgctgt tgctacctcg cccaaatctg ttttccactg 300  
ctttgacggc gagaacgtgg agagtgtctc tcctcctatg aagaaggact acaaactggc 360  
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acctgccgga ggcgtgaaga agttccagtc tgctgcttac tccaactgcg cgaagaagcc 180  
atcctgatat ccccttttggc ttctcattc tagtagttta ggatttcttt tctgacactt 240  
tgattctgac caatctctct ggctgctgc ttctgataa tcgaccagtt cccagctctt 300

gctccttgca ctctctctct catctcagca ttgtgttctg attcacctgc tccaatggct 360  
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 gttcatcttt cctggcgac agccagcccc ccacaggagc ttctctgagg aagttgcgca 180  
 cttaacctgt actttggcgg cctgaaatct ggtggtaatg cttatgtgat tggagaccag 240  
 caagacctgg acagaagtgg cacgtcttct acgccactga gtacccagag caacctgggt 300  
 taaccttgaa atgtgcatga ctggtctgga caagaagaaa gcttccgtcn ttttaaagac 360  
 taatgtgat gggaacacaa catgtgccaa ggaaatgaca aagctctctg gcactctgaa 420  
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 gccaaagccag gctattgtta tgaaaatttg tcgtctgtaa gcatgtgaac ttccgatggt 300  
 gccacatgct ggatcagtct gaataagtaa gtatgcagct ctaggtggtc agctgcgtct 360  
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 aaatgacaaa gctctctggc atctctgaaa tcatccccga gatggagatc tgcgattttg 240  
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 atgtgacgcc cgaggacggg ttcagctatg ccagttatga gggttatgggc ttggatgcc 360  
 ctgctctgtc ttatgggtgac cttgtcaaga gggctcttcg gtgctttggc ccctcggagt 420  
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 cttggtgcan aggtctatga c 501

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<210> 535  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 ggacgcgtgg gctgaaatct ggtggtaatg cttatgtgat tggagatcca gcaagaccgg 120  
 acagaagtgg cacgtcttct acgccactga gtaccagag caaccaatgg ttaacctgag 180  
 atgtgcatga ctggctctgga caagaagaaa gcttgtgtct ttttcaagac taatgcgatg 240  
 ggaacacaac atgtgccaag gaaatgacaa agctctctgg catctctgaa atcatccccga 300  
 tatggagatc tgcgattttg acttcgaacc ctgtggctac tccatgaatg cgatcatggc 360  
 tctgcattct ccacaatcca tgtgacgccc gaggacgggt tcaagctacg ccattacgan 420  
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<210> 536  
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 <212> nucleic acid

<213> Zea mays

<400> 536

gcacgactgg tctggacaag aagaaagctt gtgtcttttt caagactaat gctgatggga 60  
acacancatg tgccaaggaa atgacaaagc tctctggcat ctctgaaatc atccccgaga 120  
tgagatctg cgattttgac ttcgaaccct gtggctactc catgaatgcg atccatggct 180  
ctgcattctc cacaatccat gtgacgcccg aggacggttt cagctacgcc agttacgagg 240  
ttatgggctt ggatgccact gctctgtctt atgggtgacct tgtcaagagg gtccctccggt 300  
gcttttgccc ctcagagttt tccgttgccg tgaccatctt cggcgggctg ggccatgccg 360  
ggacatgggg aaaggcactt ggtgcanaag tctatgactg cacancatgg tggaacanga 420  
acttgccctg agcggtttct ct 442

<210> 537

<211> 421

<212> nucleic acid

<213> Zea mays

<400> 537

ctgaatagnt gtctatgcc a ttgctgctg tgaaggncn cnntgggacg ttcattcttn 60  
ctggcgacac gccagcccc cacaggagct tctctgagga agttgctgca cttaaccgct 120  
actttggcgg cctgaaatct ggtggtaatg cttatgtgat tggagatcca gcaagacctg 180  
gacagaagtg gcacgtcttc tacgccactg agtaccaga gcaaccaatg gttaaccttg 240  
aatgtgcat gactggctctg gacaagaaga aagcttctgt ctttttcaag actaatgctg 300  
atgggaacac aacatgtgcc aaggaaatga caaagctctc tggcatctct gaaatcatcc 360  
ccgagatgga gatctgcgat ttgactttg aaccctgtgg ctacttcatg aatgccaanc 420  
c 421

<210> 538

<211> 410

<212> nucleic acid

<213> Zea mays

<400> 538

gtctttttca agactaatgc tgatgggaac acaacatgtg ccaaggaaat gacaaagctc 60

tctggcatct ctgaaatcat ccccgagatg gagatctgcg attttgactt tgaaccctgt 120  
ggctactcca tgaatgcat ccatggctct gcattctcca caatccatgt gacgcccag 180  
gacggtttca gctatgccag ttatgaggtt atgggcttgg atgccactgc tctgtttat 240  
ggtgaccttg tcaagagggg ccttcgggtgc tttggccctt cggagttttc cgttgccgtg 300  
accatcttcg gggggcgtgg ccatgccggg acatggggaa aggcacttgg tgcagaggtc 360  
tatgactgca acaacatggg ggagcangag ctgcctggag gcgggcttct 410

<210> 539  
<211> 475  
<212> nucleic acid  
<213> Zea mays

<400> 539

gnggnnnnnn gnttnttagc ngcngcacc aatcaaagan tcgcgggtcg anncacgcgc 60  
cnctccctcn atctccagca ttgtgttctg attcacctgc tccaatggct gttctttcgc 120  
tgctgatgct tccccggtct cagctatcgg gtttgagggc tatgagaagc gccttgaatc 180  
acattctctg aggcacctgt ctttgtggac cctcatgggc gtggtttgcg tgcccttcca 240  
gggccagat tgactctgtt ctggatcttg cacggtgcac aattgtgtct gagcttccaa 300  
caaggatttc gactcatatg tcctttctga gtcaagcttg tttatctatc ctctaagatt 360  
gtcatcaaga cctgtggcac taccaagctc ctgctacca ttccaagaat cctgagcttg 420  
ctgaagaact gtctatgcca cttgctgcgt gaaataatcc cgttggaac gtcac 475

<210> 540  
<211> 500  
<212> nucleic acid  
<213> Zea mays

<400> 540

ggnnnnnnnn nngttaactt cnacgccggt cggtaacanag tccnggaatt cccgggtcgc 60  
ccacgcgtcc gccacgcgt ccgcaaagct ctctggcatc tctgaaatca tccccgagt 120  
gagatctgcg attttgactt cgaaccctgt ggctactcca tgaatgcat ccatggcctg 180  
cattctccac aatccatgtg acgcccagg acggtttcag ctacgccagt tacgagttat 240  
gggcttggat gccactgctc tgtcttatgg tgacctgtc aagaggggcc tccgggcttt 300

ggccccctcag agttttccgt tgccgtgacc atcttcggcg ggcgtggcca tgccggacat 360  
 ggggaaaggc acttggtgca gaggtctatg actgcaacaa catggtggaa cagagctgcc 420  
 tggaagcggg ctctctgtgt aacaaaactt tcgtncgct gaaaancccg ttccaactcc 480  
 ccnaaatcnt ttttcaanng 500

<210> 541  
 <211> 508  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 541

agnnaggagn ggtgnnncat ngntagngg ntnaannaaa tncncgtacc ggtccggaat 60  
 cccgggtcga cccacgcgtc cgctccagca ttgtgttctg attcacctgc tccaatggct 120  
 gttctttctg ctgctgatgn ttccccggtc tcagctatcg ggtttgaggg ctatgagaag 180  
 cgccttgaga tcacattctc tgaggcacct gtctttgtgg accctcatgg gcgtgggttg 240  
 cgtgccctct ccagggccca gattgactct gttctggatc ttgcacggtg cacaattgtg 300  
 tccgagctct ccaacaagga ttctgactca tatgtcctct ctgagtcaag cttgtttatc 360  
 tctctctga agattgtcat caagacctgt ggcactacca agctcctgct caccattcca 420  
 agaatccttg agcttgctga agaagctgtc tatgccactt gctgctgtga aagtacttcc 480  
 gtgggacgtc atctttcctg gtgcacac 508

<210> 542  
 <211> 468  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 542

ggnnnnngta ctttanntnn cagaggcctg gtcaagnnca agcgggtcga nncangcntn 60  
 nncagactt ctgcgntctc agctatcggg tttagggct atgagaagcg ccttgagaca 120  
 cattctctga ggcacctgtc tttgtggacc ctcatggcg tggtttgcgt gccctctcag 180  
 ggcccagatt gactctgttc tggatcttgc acggtgcaca attgtgtctg agctctcaac 240  
 aaggatttcg actcatatgt cttttctgaa gtcaagcttg tttatctatc ctctgagatt 300  
 gtcatcaaga cctgtggcac taccaagctc ctgctacca ttccaagaat ccttagcttg 360



ctgaagagct gtctatgcc a cttgctgctg tgaagtactc ccgtgggaacg ttctctttcc 420  
 tgggcgcaca gccagccccc cacangagct tctctganga agttgctg 468

<210> 543  
 <211> 494  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 543

ggnngnaact tctacgcgc cggtttcggt cgncaaattc ccgggtaaac ccacgcgctg 60  
 nccacgcgtc cgcccacgcg tccgggggtcc ttccggtgctt tggccctcgc gagttttcgt 120  
 tgccgtgacc atcttcggcg ggcgtggcca tgccgggaca tggggaaagg cacttgggca 180  
 gaggtctatg actgcaacaa catggtggag caggagctgc ctggaggcgg gctcctgtgt 240  
 accagagctt ctgtgctgct gaagacgctg ttgctacctc gcccaaactc gttttcactg 300  
 ctttgacggc gagaacgtgg agagtgtcc tctcctatg aagaaggact acaactggct 360  
 aatcttctat gctgggagga ggaagcggat gccatggagg agaangcggg agtcttgatg 420  
 agtaagacgg gcttctgggg tccaattgcc tccganttg ttaatttaaa actcccaaact 480  
 ttntgggttg cctt 494

<210> 544  
 <211> 511  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 544

agtgtnaann ggntggnnng nnangttntn ctgtacngaa nngnanntc gaccnanncg 60  
 tacgctttcg gctacgnnat tctantagtt taggatttct tttctgacac tttgattctg 120  
 accaatctcn ctggcctgct gcttctgat aatcgaccag ttccccagtc ttgctccttg 180  
 cactcctccc tccatctcca gcattgtgtt ctgattcacc tgctccaatg gctgntcttt 240  
 ctgctgctga tgcttcccg tctcagctat cgggtttgag ggctatgaga agcgccttga 300  
 gatcacattc tctgaggcac ctgtctttgt ggacctcat ggcgtgggtt tgctgcct 360  
 ntncaggggc cagattgact ctggtctgga tcttgacagg tgcacaattg tgtncgagct 420  
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naaaatggca atcaaaaacc tgtgggacta c

511

<210> 545  
<211> 478  
<212> nucleic acid  
<213> Zea mays  
  
<400> 545

ttnnntgceen ngccgggtcna ngtcacaang ncccgggtcg acccacgcgt ccnttggggt 60  
cacctgctcc aatggctgtt ctttctgctg ctgatgttc cccggtctca agctatcggt 120  
ttgagggcta tgagaagcgc cttgagatca cattctctga ggcacctgtc tttgtggccc 180  
tcatgggctg ggtttgctg ccctctccag ggcccagatt gactctgttc tggatctgca 240  
cgggtgcacaa ttgtgtctga gctctccaac aaggatttcg actcatatgt ccttttgagt 300  
caagcttggt tatctatcct ctgaagattg tcatcaagac ctgtggcact accagctcct 360  
gtcaccatt ccaagaatcc ttgagcttgc tgaagaactg tctaagccac ttgtgctgtg 420  
aaataccceen tgggaaattn aaacctttcc ngngcaaan caacccccca cagagctt 478

<210> 546  
<211> 533  
<212> nucleic acid  
<213> Zea mays  
  
<400> 546

ggnnaggnc a ggtttgann nggnatggg ggaaatnntt cgcccgaat tcccgggtcg 60  
accacgcgt cgtgaagat tgtcatcaag acctgtggca ctaccaagct cctgctcaca 120  
attccaagga tcttagagct tgctgaagag ctgtctatgc ctcttgctgc tgtgaagtac 180  
tcccgaggga cgttcatctt tcttggcgca cagccagccc cccaccggag cttctccgag 240  
gaagttgctg tacttaaccg atactttggg ggctgaagt ctggtggcaa tgcttatgtg 300  
attggagatg cagcaagacc aggacagaag tggcacatct actacgccac tgagtacca 360  
nagcaacca tggtaacct tgagatgtgc atgactggtc tggacacgaa gaaagcttca 420  
gtcttcttca agactaatgc tgatggcaac acaacatgtt ccaagggaaat gaccaaactt 480  
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<210> 547

<211> 485  
 <212> nucleic acid  
 <213> Zea mays

<400> 547

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 aggactacaa gctgggctaatt cttctctgct gggaggagga agcggatgcc atggagggaa 180  
 ggcggggagtg cttgatgagt aagacgggct tctggggctcg atttgcttct gagttgttat 240  
 tttatatcgt cgcaatttcg tgggtgtcgt ttggttatcc tgtgaagcag ccaagcaggc 300  
 tattgttatg aaaatttgct gtctgtaagc atgtgaactt ccgatgttgc cacagctgga 360  
 tnagtctgaa taagtaagta tgcaactcta ngtggtcanc tgcgtctaac acatgaagca 420  
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 aattt 485

<210> 548  
 <211> 442  
 <212> nucleic acid  
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<400> 548

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 ctgctctgtc ttatgggtgac cttgtcaaga gggtccttcg gtgctttggc cctcggagt 180  
 tttccgttgc cgtgaccatc ttcggcgggc gtggccatgc cgggacatgg ggaaaggcac 240  
 ttggtgcaaa ggtctatgac tgcaacaaca tgggtggaaca agagctgcct ggaggcgggc 300  
 tectcgtgta ccaaancctc tgtgctgctg aagacgctgt tgctacctcc ccaaactctgt 360  
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 caaactggct aatcttntat gg 442

<210> 549  
 <211> 449  
 <212> nucleic acid  
 <213> Zea mays

<400> 549

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tgnntggaga tccagcaaga cctggacaga aatggcacgt cttctacgcc actgagtacc 120

cagagcaacc aatgggtaac cttgaaatgt gcatgactgg tctggacaag aagaaagctt 180

ctgtcttttt caagactaat gctgatggga acacaacatg tgccaaggaa atgacaaagc 240

tctctggcat ctctgaaatc atccccgaga tggagatctg cgattttgac tttgaaccct 300

gtggctactc catgaatgcg atccatggct ctgcattctc cacaatccat gtgacgcccg 360

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ngggancctt tcnagaaggg tccttcggg 449

<210> 550

<211> 512

<212> nucleic acid

<213> Zea mays

<400> 550

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tcctganant cnatcatgtt cncagggtct tgctccttgc actcctccct ccattctcagc 180

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ttctggatct gcaengtgca cnnttggtc tgagctctcc aacacggnnt tegtcaata 420

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<211> 448

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<213> Zea mays

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ctgcacttaa ccgctacttt ggcggcctga aatctggtgg taatgcttat gtgattggag 240  
atccagcaag acctggacag aagtggcacg tcttctacgc cactgagtac ccagagcaac 300  
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ccacngatcc atgtgacgcc tgaggatggg ttcagctacg ccagttatga ggttatgggc 180  
ttggatgcca ctgccctgtc ttacggtgac cttgtcaaga gggtccttgg gtgcttcggc 240  
ccctcagaat tttctgtcgc cgtgaccatc ttcggcgggc ggngccaagc tgggacatgg 300

ggaaaggaac ttggtgcgga ggcttatgac tgcaacaaca tggtcgagca ggagctgcct 360  
ggaggtggga tcctcatcta ccaaagcttc ttgtgctgnt gaaanacccc gttgctagct 420  
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<210> 554  
<211> 347  
<212> nucleic acid  
<213> Zea mays

<400> 554

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tgctttggcc cctcagagtt ttccgttgcc gtgaccatct tcggcgggag tggccatgcc 180  
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gagctgcctg gaggcgggct cctcgtgtac cagagcttct gtgctgctga agacgctgtt 300  
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<210> 555  
<211> 435  
<212> nucleic acid  
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<400> 555

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tggagatctg cgattntgac ttccnaccct gtggctactc catgaatgcg atccatggct 180  
ctgcattctc cacaatccat gtgacgcccg agnccggttt cagctacncc agttacgagg 240  
ttatgggctt ggatgccact gntntgtctt atggcgacct tgtcaagagg gtccttcggt 300  
gctttgcccc tcagagtttt acgangccgt gaccatcttc ngcngaagtg nccatgcctg 360  
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agcngccttg angca 435

<210> 556  
<211> 334  
<212> nucleic acid

<213> Zea mays

<400> 556

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cgctgctgaa gacgctgttg ctacctcgcc caaatctgtt ttccactgct ttgacggcga 120  
gaacgtggag agtgctcctc ctctatgaa gaaggactac aaactagcta atcttctctg 180  
ctgggaggag gaagcggatg ccatggagga gaaggcggga gtgcttgatg agtaagacgg 240  
gttctctggtg tcgatttgct tctgagttgt ttattttata tcgtcgcaat ttcgtgggtg 300  
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<210> 557

<211> 404

<212> nucleic acid

<213> Zea mays

<400> 557

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tgaagattgt catcaagacc tgtggcacta ccaagctcct gctcaccatt ccaagaatcc 180  
ttgagcttgc tgaagagctg tctatgccac ttgctgctgt gaagtactcc cgtgggacgt 240  
tcatctttcc tggcgcacag ccagccccc acaggagctt ctctgaggaa gttgctgcac 300  
ttaaccgcta ctttggcggc ctgaaatctg gtggtaatgc ttatgtgatt ggagatccag 360  
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<210> 558

<211> 430

<212> nucleic acid

<213> Zea mays

<400> 558

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gatcttgcac ggtgcacaat tgtgtccgag ctctncaaca aggatttcga ctcatatgtc 180  
ctttctgagt caagcttggtt tatctatcct ctgaagattg tcatcaagac ctgtggcact 240

accaagctcc tgctcaccat tccaagaatc cttgagcttg ctgaagaact gtctatgcca 300  
 cttgtctgntg tgaaagtact cccgtgggga cgttcatctt tcctggcgca caagncaggn 360  
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 ccttgaaant 430

<210> 559  
 <211> 319  
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 gatggagatc tgcgattttg acttogaacc ctgtggctac tccatgaatg cgatccatgg 180  
 ctctgcattc tocacaatcc atgtgacgcc cgaggacggg ttcaagctacg ccagttacga 240  
 ggttatgggc ttggatgcca ctgctctgtc ttatgggtgac cttgtcaaga gggtcctccg 300  
 gtgctttggc ccctcagag 319

<210> 560  
 <211> 346  
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 ctttttcaag actaatgctg atgggaacac aacatgtgcc aaggaaatga caaagctctc 180  
 tggcatctct gaaatcatcn ttnagatgga gatctgcatg tttgacttcg aaccctgtgg 240  
 ctactccatg aatgcgatcc atggctctgc attctccaca atccatgtga cgcccaggga 300  
 cggtttcagt acgccattac gaggttatgg gcttggatgc catgct 346

<210> 561  
 <211> 317  
 <212> nucleic acid  
 <213> Zea mays



<400> 561

gtacccagag caaccaatgg ttaaccttga gatgtgcatg actggtctgg acaagaagaa 60  
agcttgtgtc tttttcaaga ctaatgctga tgggaacaca acatgtgcc aggaaatgac 120  
aaagctctct ggcctctctg aaatcatccc cgagatggag atctgogatt ttgacttcga 180  
accctgtggc tactccatga atgcgatcca tggctctgca ttctccacaa tccatgtgac 240  
gcccaggagc ggtttcagct acgccagtta cgaggttatg ggcttgatg ccactgctct 300  
gtcttatggg gaccttg 317

<210> 562

<211> 342

<212> nucleic acid

<213> Zea mays

<400> 562

ccgaggaggt ttcagctatg ccagttacga ggttatgggc ttggatgcc ctgctctatc 60  
ttatggtgac cttgtcaaga gggtcctccg gtgctttggc ccctcggagt tttcogttgc 120  
cgtgaccatc ttcggcgggc gtggccatgc cgggacatgg ggaaaggcac taggtgcaga 180  
ggcttatgac tgcaacaaca tgggtggagca ggagctgcct ggaggcgggc tctcgtata 240  
ccagagcttc tgcgctgctg aagacgctgt tgctacctg cccaaatctg ttttccactg 300  
ctttgacggc gagaacgtgg agagtgtcct cctcctatga ag 342

<210> 563

<211> 314

<212> nucleic acid

<213> Zea mays

<400> 563

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agtagtttag gattttotitt ctgacacttt gattctgacc aatctctctg gctgctgct 120  
tctgataat cgaccagttc ccagctcttg ctcttgacac tctccctcc atctccagca 180  
ttgtgttctg attcaactgc tccaatggct gttctttctg ctgctgatgc tccccggtc 240  
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<210> 564  
 <211> 541  
 <212> nucleic acid  
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 <400> 564

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 atctgttttc cactgctttg acggcgagaa cgtggagagt gtcctcctc ctatgaagaa 240  
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 ccgtgaccat cttcggcggg cgtggccatg ccgggacatg gggaaaggca cttggtgcag 180  
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<400> 618

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 tctagtagtt taggatttct tttctgacac tttgattctg accaatctct ctgggcctgc 180  
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<210> 654  
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 <400> 654  
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 caacaacatg gtggagcang agctgcctgg aggcgggctc ctcntgtacc aaagctttgt 180  
 nctgntgaag acgcngttgc tanctcgccc aaatccgttt tccantgntt tgacnggaaa 240  
 acnttganaa tntcctcct cctangaaga aagantacaa actggntaat cttctanctg 300  
 ggangaggaa acngattccn nggnngagaa agcgggagtg cctgatnngt attaaggntt 360  
 ctggggncna ttttcgtctn agttngtann ttttaagcnt cccaanattc cccgggtgtc 420  
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 nccgnnng 488

<210> 655  
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 <212> nucleic acid  
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<400> 655  
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 cttgtttatc tatcctctga agattgtcat caagacctgt ggccactacca agctcctgct 180  
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<400> 656  
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 gagatgcagc aagaccagga cagaagtggc acatctacta cgccactgag taccagagc 180  
 aaccaatggt caaccttgag atgtgcatga ctggctctga cacgaagaaa gcttcagtct 240  
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<210> 657

<211> 250  
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 gaagagctgt ctatgccact tgctgctgtg aagtactccc gtgggacgtt catctttcct 120  
 ggcgcacagc cagcccccca caggagcttc tctgaggaag ttgctgcact taaccgctac 180  
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 cagaagtggc 250

<210> 658  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 658  
  
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 atgtcctttc tgagtcaagc ttgtttatct atcctctgaa gattgtcatc aagacctgtg 180  
 gcactaccaa gctcctgctc accattccaa gaatccttga gcttgctgaa gagctgtcta 240  
 tgccacttgc tgctgtgaag t 261

<210> 659  
 <211> 261  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 659  
  
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 gttaaccttg agatgtgcat gactggctctg gacaagaaga aagcttgtgt ctttttcaag 120  
 actaatgctg atgggaacac aacatgtgcc aaggaaatga caaagctctc tggcatctct 180  
 gaatcatccc ngagatggag atctgcgatt tngacttcga accctgnggc tactccatga 240  
 ntgcgatcca tggctctgca t 261

<210> 660

<211> 454  
 <212> nucleic acid  
 <213> Zea mays

<400> 660

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gcgtctggtg gtggtgaagg ggagaaattc gtgagatctg ttccggatca ggcgtgcgag  180
ctcgggaatc aggggtttca cacatagctt cgtcgatttg aatttgatgt actaatggag  240
tctaaggggtg gcaaaaagtc tagcagtagt cgttccatga tgtatgaagc tccccttggc  300
tacagcattg aggacgttcg acctgccgga ggcgtgaaga agttccagtc tgctgcttac  360
tccaactgcg cgaagaagcc atcctgatat cccttttggc ttccctcattc tagtaagttt  420
aggattttctt ttctgacact ttgattctga ccca                                454

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<210> 661  
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 <212> nucleic acid  
 <213> Zea mays

<400> 661

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cacgtcttct acgccactga gtaccagag caaccaatgg ttaaccttga gatgtgcatg  180
actggtctgg acaagaagaa agcttgtgtc tttttcaaga ctaatgctga tgggaacaca  240
acatgtgc                                         248

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<210> 662  
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 <212> nucleic acid  
 <213> Zea mays

<400> 662

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atggctctgc gttctccacg atccatgtga cgctgagga tgggttcagc tacgccagtt  120
atgaggttat gggcttggat gccactgccc tgtcttacgg tgaccttgtc aagagggtcc  180

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ttgggtgctt cggccctcg gaattttctg tcgccgtgac catcttcggc gggcggggcc 240  
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<212> nucleic acid  
<213> Zea mays  
<400> 663

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ggagagtgtc cctcctccta tgaagaagga ctacaagctg gctaattctt tctgctggga 180  
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ggggtcg 247

<210> 664  
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tggacagaag tggcacgtct tctacgccac tgagtaccca gagcaaccaa tggttaacct 180  
tgagatgtgc atgactggtc tggacaagaa gnnagcttgt ggtcttttca agactaatgc 240  
tgatgggaac acaa 254

<210> 665  
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<400> 665

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tcttcggcgg gcgtggccat gccgggacat ggggaaaggc acttggtgca gaggtctatg 180

actgcaacaa catggtggag caggagctgc ctggaggcgg gtcctcgtg taccagagct 240  
tctgtgctgc tga 253

<210> 666  
<211> 248  
<212> nucleic acid  
<213> Zea mays  
  
<400> 666

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gtaatgctta tgtgattgga gatccagcaa gacctggaca gaagtggcac gtcttctacg 180  
ccactgagta cccagagcaa ccaatgggta accttgagat gtgcatgact ggtctggaca 240  
agaagaaa 248

<210> 667  
<211> 291  
<212> nucleic acid  
<213> Zea mays  
  
<400> 667

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ctttgtggac cctcatgggc gtggtttgcn tgccctctcc agggcccaga ttgactctgt 180  
tctggatctt gcacgggtgca caattgtgtc tgagctctcc aacaaggatt cgatcatagt 240  
ccttinctgan tnagcttggt taactatcct ctgaagatgt catcaagact g 291

<210> 668  
<211> 247  
<212> nucleic acid  
<213> Zea mays  
  
<400> 668

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ngagtacca gagcaaccaa tggttaacct tgagatgtgc atgactggtc tggacaagaa 120  
gaaagcttgt gtctttttca agactaatgc tgatgggaac acaacatgtg ccaaggaaat 180

gacaaagctc tctggcatct ctgaaatcat ccccgagatg gagatctgcg attttgactt 240  
cgaaccc 247

<210> 669  
<211> 445  
<212> nucleic acid  
<213> Zea mays  
  
<400> 669

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gagttctagg tagagaatcg gntttacttg tctcagcccg gggctctgctg cgtctgggtg 120  
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caaaaagtct agcagtagtc gttctatgat gtatgaagct ccccttggtc acagcattga 300  
ggacnttcga cctgcncgag gcgtgaaaaa nttccagtct gntgcttact ncaactgnnc 360  
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ctgacacttt gattctgacc aatct 445

<210> 670  
<211> 276  
<212> nucleic acid  
<213> Zea mays  
  
<400> 670

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gtcttggtga cctcatggg cgtgggttgn gtgcnctctc cagggncag attgactctg 180  
ttctggatct tgcacggtgc acaatgtgtc cgagctctcc aacaaggatt tcgactcana 240  
tgtctctctc tgagtcaagc ntgtntatct atccnc 276

<210> 671  
<211> 283  
<212> nucleic acid  
<213> Zea mays  
  
<400> 671

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 ccaactgogc aaagaagcca tcttgatata ccttttggnt tcctcgttct agtagtttag 180  
 gattttcttt ctgacacttc gnttcttacc aatccccctg gcctgctgct ttcttgacaa 240  
 togaccagtt cccagtcctt gctccctgca gtctccctc ctc 283

<210> 672  
 <211> 284  
 <212> nucleic acid  
 <213> Zea mays

<400> 672

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 gctccaatgg ctgttctttc tgctgctgat gttccccggt ctgagctatc gggtttgagg 180  
 gctatgagaa ggccttgag atcacattct ctgaggcacc tgtctttgtg gaccctcatg 240  
 ggcggtggtt gcgtgccctc tcaagggccc agattgactc tggt 284

<210> 673  
 <211> 443  
 <212> nucleic acid  
 <213> Zea mays

<400> 673

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 tggttattct gngaagcanc caagccaggc tattgntatg aaaatttgct gtntgtaagc 180  
 atgtgaactt ncgatgttgc cacatgctgg atcagtctga ataagtaagt atgcagctct 240  
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 ctttttggnt atgaattaca annngctttt tngaaaaaan aatactggcg gnnntntgttc 360  
 aanaaatggt tttgagcttt nctttngta tcttttttga agaaccggtc taattgacgc 420  
 tttaaattcnc atggagtgggt naa 443

<210> 674



<211> 254  
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 <213> Zea mays  
  
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 tcgtgtacca gagcttctgt gctgctgaag acgctgttgc tacctcgccc aaatctgttt 180  
 tccactgctt tgacggcgag aacgtggaga gtgctcctcc tcctatgaag aaggatacaa 240  
 gctggctaatt cttc 254

<210> 675  
 <211> 272  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 675  
  
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 actggtctgg acacgaagaa agcttcagtc ttcttcaaga ctaatgctga tggcaacaca 180  
 acatgtgcc aaggaaatgac gaagctctct ggtatctctg aaattatccc tgagatggag 240  
 atctgtgatt ttgacttcga accctgcggc ta 272

<210> 676  
 <211> 240  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 676  
  
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 caatggttaa ccttgagatg tgcattgactg gtctggacaa gaagaaagct tgtgtctttt 120  
 tcaagactaa tgctgatggg aacacaacat gtgccaaagga aatgacaaag ctctctggca 180  
 tctctgaaat catccccgag atggagatct gcgattttga cttcgaaccc tgtggctact 240

<210> 677  
 <211> 243  
 <212> nucleic acid

<213> Zea mays

<400> 677

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ggtaatgctt atgtgattgg agatccagca agacctggac agaagtggca cgtcttctac 180  
gccactgagt acccagagca accaatgggt aaccttgaga tgtgcatgac tggctctggac 240  
aag 243

<210> 678

<211> 243

<212> nucleic acid

<213> Zea mays

<400> 678

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tatgactgca acaacatggt ggagcaggag ctgcctggag gcgggctcct cgtgtaccag 180  
agcttctgtg ctgctgaaga cgctgttgct acctcgccca aatctgtttt cactgcttt 240  
gac 243

<210> 679

<211> 251

<212> nucleic acid

<213> Zea mays

<400> 679

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ggacgttcga cctgccggag ggtgaagaag ttccagtctg ctgcttactc caactgcgcg 180  
aagaagccat cctgatatcc cttttggcct cctcattcta gtagtttagg atttcttttc 240  
tgacactttg a 251

<210> 680

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 680

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cgaggaagtt gctgtactta accgatactt tggggggcctg aagtctgggtg gcaatgctta 180  
tgtgattgga gatgcagcaa gaccaggaca gaagtggcac atctactacg cactgagta 240  
cccagagcaa ccaatggtca accttgag 268

<210> 681

<211> 249

<212> nucleic acid

<213> Zea mays

<400> 681

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atccatggct ctgcattctc cacaatccat gtgacgccg aggacggttt cagcnatgcc 180  
agttatgagg ttatgggctt ggatgccact gctctgtctt atggtgacct tgtcaagagg 240  
gtccttcgg 249

<210> 682

<211> 241

<212> nucleic acid

<213> Zea mays

<400> 682

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atctgggtggt aatgcttatg tgattggaga tccagcaaga cctggacaga agtggcacgt 180  
cttctacgcc actgagtacc cagagcaacc aatgggtaac cttgagatgt gcatgactgg 240  
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<210> 683

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 683

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tggctactcc atgaatgcga tccatggctc tgcattctcc acaatccatg tgacgcccga 180  
ggacggtttc agctacgccg gttacgaggt tatgggctnt anttgccatg ctctgtctta 240  
tggtgacctt gtcaagaggt cctccggt 268

<210> 684

<211> 264

<212> nucleic acid

<213> Zea mays

<400> 684

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gggagtgttt gatgagtaag acgggcttct ggtgtcgatt tgcttctgag ttgtttatTT 180  
tatatcgtcg cnatttcgtg gttgtcgttt ggttatctgt gaagcagcca agccaggcta 240  
ttgttatgaa aattgtcgtc tgta 264

<210> 685

<211> 268

<212> nucleic acid

<213> Zea mays

<400> 685

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aagactaatg ctgatgggaa cacaacatgt gccaaaggant ganaancctc tctggcatct 180  
ctgaaatcat cccccgagatg gagatctgcg attttgactt cgaaccctgt ggctactcca 240  
tgaatgcgat ccatggctct gcattctc 268

<210> 686

<211> 332

<212> nucleic acid

<213> Zea mays

<400> 686

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gggagtgctt gatgagtaan acgggcttct ggggtcgatt tgcttctgag ttgtttatct 180  
tatatcgctg caatttcgtg gttgtcgctt ngttattctg tgaagcagcc aagccaagct 240  
attgttatga nntttgtcgt ctgtaagcat gtgacttccg ccgttgccac atgctggatc 300  
agtctgaata gtaagtngca gctctagtgg nc 332

<210> 687

<211> 237

<212> nucleic acid

<213> Zea mays

<400> 687

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attctgtgaa gcagccaagc caggctattg ttatgaaaat ttgtcgtctg taagcatgtg 180  
aacttcgat gttgccacat gctggatcag tctgaataag taagtatgca gctctag 237

<210> 688

<211> 286

<212> nucleic acid

<213> Zea mays

<400> 688

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cccggtctca gctatcgggt ttgagggcta tgagaagcgc cttgagatca cattctctga 120  
ggcacctgtc ttgttgacc ctcatggcgt ggtttgctg ccctctccag ggcccagatt 180  
gactctgttc tggatcttgc acggtgcaca atgtgtccga gctctccaac aaggatttcg 240  
actcatagtc ctctctgagt caagcttggtt tatctatctc tgaaga 286

<210> 689

<211> 300

<212> nucleic acid

<213> Zea mays

<400> 689

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tccccttggc tacagcnttg aggacgttcg acctgccga cggatgaagaa gttccagtct 180  
gctgcttact ccaagtgcgc gangaagcca tcttgatctt cttttggctt cctcattct 240  
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<210> 690

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 690

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ggggggctgg ctgtgcacca ggaaagatga acgtcccang ggagtacttc acagcagcna 120  
gtggcataga cagctcttca gcaagctcaa ggattcttgg aatggtgagc aggacttggt 180  
agtgccacag gtcttgatga caatcttcag aggatagatn aacaagcttg actcatagag 240  
gacatatgat ccgaaatcct tgtggagagc tcgacaca 278

<210> 691

<211> 525

<212> nucleic acid

<213> Zea mays

<400> 691

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tctcagcccg gggctctgtg cgtctgggtg tggatgaagg gagaaattcg tgagatctgt 240  
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gttccagtct gctgcttact ccaactgcgc gaagaaccat cctgatatcc ctttttgctt 480  
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<213> Zea mays

<400> 692

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<212> nucleic acid  
<213> Zea mays

<400> 693

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<400> 694

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ntncccgnga tggannntgc gnt

263

<210> 695  
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<400> 695

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atctg 245

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<400> 696

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tccagggcc agattgactc tgttctggat cttgcacggt gcacaattgt gtctgagctc 180

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<400> 697

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tctgtgctgc tgaagacgct gttgctacct cgcccaaadc tgtttccac tgctttgacg 180

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 cgtggagagt gctcctcctc ctatgaagaa ggactacaag ctggctaate ttctctgctg 180  
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 ctgagaggtg ccgggaccga ggaactggcc gagttctagg tagagaatcg gttttgcttg 180  
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 tccggatcag gcgtgcgagc tcgggaatca ggggtttcac acatagcttc gtcgatttga 300  
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 gtatgaagct ccccttggtc acagcattga ggacgttcga cctgccggag gcgtgaagaa 420

gttccagtct gctgcttact ccaactgcg c gaagaagcca tcttgatata ccttttggct 480  
tcttcattct agtagtt 497

<210> 701  
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<212> nucleic acid  
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<400> 701

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tacagcattg aggacgttcg acctgccgga ggggaagaa gttccagtct gctgcttact 180  
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<210> 702  
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<400> 702

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tgctgcttac tccaactgcg cgaagaagcc atcctgatan ccttttnggc ttcctcattn 240  
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<210> 703  
<211> 225  
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<400> 703

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ttctggatct tgcaagggtgc acaattgtgt ccgagctctc caacaaggat ttcgactcat 180

atgtcctctc tgagtcaagc ttgtttatct atcctctgaa gattg 225

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 <211> 289  
 <212> nucleic acid  
 <213> Zea mays

<400> 704

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 gtttgagggc tatgagaagc gccttgagat cacattctct ganncacctg tctttgtgga 180  
 ccccatggc agcgggttgc gtgccctctc caggtccag attgactctg ttctggatct 240  
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<210> 705  
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 <213> Zea mays

<400> 705

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<210> 706  
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<400> 706

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 ctcattctag tagtttagga tttcttttct gacactttga ttctgaccaa tctctctggc 180  
 ctgctgcttc ctgataatcg accagttccc cagtcttgc c 221

<210> 707  
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 <212> nucleic acid  
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 cncanggcgg ggg 253

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<210> 710  
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<400> 710

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gaaaggaatt gg 252

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<212> nucleic acid  
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<400> 713

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<400> 715

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<210> 721  
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<210> 722  
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 tttgtggacc cccatggcag cggtttgcgt gccctctcca ggtcccagat tgactctgtt 240  
 ctggatcttg cacggtgcac 260

<210> 726  
 <211> 536  
 <212> nucleic acid

<213> Zea mays

<400> 726

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cgnatgngac cgacgcgtcc gcaangaatt ccctccccta gccgcgcgt ccgtcgctcg 120  
tctccctgag aggtgccggg ncnggggaac tggccgagta ctaggtagag aatcggtttt 180  
tcttgtctca gcccggggtc tgctgcgtct ggtggtggtg aaggggagaa attcgtgaga 240  
tctgttccgg atcaggcgtg cgagctcggg aatcangggg ttcacacata gcttcgtcga 300  
tttgaatttg atgtactaat ggagtctaag ggtggcaaaa agtctagcag tagtcgttct 360  
atgatgtatg aagctnccct tggctacagc attgaggacc gttcgacctg ccggangcgt 420  
gaagaagttc cagtctnctg cttacttcaa ctgccnaag aagccatcct natannccct 480  
ttggctttct tattctaata gntttangat ttctttttct ganaccnnt attttt 536

<210> 727

<211> 456

<212> nucleic acid

<213> Zea mays

<400> 727

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gcgtctggtg gtggtgaagg ggagaaattc gtgagatctg ttccggatca ggcggtcgag 180  
ctcggaatc aggggtttca cacatagctt cgctgatttg aatttgatgt actaatggag 240  
tctaagggtg gcaaaaagtc tagcagtagt cgttctatga tgtatgaagc tccccttggc 300  
tacagcattg angacgttcg acctgccnga ngcgtgaaga aattncaagt ctgctgctta 360  
ctccaactgc gccaaaaaac catnctgata ttcttttggg ctttcctaata ctagtaagtt 420  
aaggatttct tttctgacac ttttgaatct gancca 456

<210> 728

<211> 196

<212> nucleic acid

<213> Zea mays

<400> 728

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 ggTtattctg tgaagcagcc aagccaggct attgttatga aaatttTgTcg tctgtaagca 120  
 tgtgaacttc cgatgttgcc acatgctgga tcagtctgaa taagtaagta tgcagctcta 180  
 ggtggTcagc tgcgtc 196

<210> 729  
 <211> 218  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 729

gattgactct gttctggatc ttgcacggTg cacgatcgTg tccgagctct ccaacaagga 60  
 ttttgactcc tatgttctct ccgagtcaag cttgttcatc tatcctctga agattgtcat 120  
 caagacctgt ggcactacca agctcctgct cacaattcca aggatcctag agcttTgctga 180  
 agagctgtct atgcctcttg ctgctgtgaa gtactccc 218

<210> 730  
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 <212> nucleic acid  
 <213> Zea mays  
 <400> 730

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 naccgntctt gtcttatggT gacctgtTca agagggtcct ccggtgcttt ggccccTcga 120  
 gttttccgTt gccgtgacca tcttcggcg gcgTggccat gccgggacat ggggaaagca 180  
 cttggTgcag aggtctatga ctgcaacaac atggTggagc aggagctgcc tggagggggc 240  
 tctcgtgta ccagagctna c 261

<210> 731  
 <211> 306  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 731

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 gtggcaaaaa gtctagcagt agtcgtctat gatgtatgaa gctccccttg gctacagcat 120

tgaggacggt cgacctgccg gagngtgaag aagttccagt ctgctgctta ctccaactgc 180  
 gcgaagaagc catcctgata tcctttggct tcctcatcta gtagtttagg attctttctg 240  
 acatttgatt ctgacaatct cctggctgtg ttctgataac gacaatcnca tctgtcttga 300  
 tctcct 306

<210> 732  
 <211> 295  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 732

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 aagaagccat cctgatatcc cttttggctt cctcgttcta gtagtttagg atttcttttc 120  
 tgacaattcg attcttacca atccccctgg cctgcngctt tcctgataat cgaccagttc 180  
 cccagtcttg ctccctgcag tcctccctcc tccatctcca gcgttggtgt ctgactcacc 240  
 tgcaccaatg gctgttcttt ctgctgctgg tgctccccgg gctnaagtat ngggt 295

<210> 733  
 <211> 532  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 733

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 cgtccgcnc a cngcgtccgc ccacgccgtc cgcccacgcg tccggggcaa atgcttatgt 120  
 gattggagat gcagcaagaa ccagggacag aagtggcaca tctactacgc cactgagacc 180  
 cagagcaacc aatggtcaac cttgagatgt gcatgactgg tctggacacg aagaaacttc 240  
 agtcttcttc aagactaatg ctgatggcaa cacaacatgt gccaaaggaaa tgacgagctc 300  
 tctggttaanc tctgaaatta tcctgagat ggagatctgt gatcttgact tcgaaccctg 360  
 cggtactcc catgaatgca atccatgggc tctgcgttct ccacgattca tgtgacgcct 420  
 gaggatgggt tcaagctacg ccaagttatg aagttatggg gcntggatgc catggncctg 480  
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<210> 734

<211> 269  
 <212> nucleic acid  
 <213> Zea mays  
  
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 tctccctcc tccatctcca gcgttggtt ctgactcacc tgcaccaatg gctgttcttt 120  
 ctgctgctgg tgctccccg gcctcagcta tcgggtttga gggctatgag aagcgcttg 180  
 agatcacatt ctctgaggca cctgtctttg tggaccccca tggcagcggg ttgcgtgccc 240  
 tctccaggtc ccagattgac tctgttctg 269

<210> 735  
 <211> 231  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 735  
  
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 tagtcgttcc atgatgtatg aagctcccct tggctacagc attgaggacg ttcgacctgc 120  
 cggagggtga agaagttcca gtctgctgct tactccaact gcgcgaagaa gccatcctga 180  
 tatccctttt ggcttccgca tnnaggagtt agggtttcta cccnnccngn g 231

<210> 736  
 <211> 220  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 736  
  
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 aagcatgtga acttccgatg ttgccacatg ctggatcagt ctgaataagt aagtatgcag 120  
 ctctaggtgg tcagctgcgt ctaccacaat gagcatgaac gtatggagaa atatctgtga 180  
 acccattttg tttatgaata agatttggtt ttttcgagtt 220

<210> 737  
 <211> 442  
 <212> nucleic acid  
 <213> Zea mays

<400> 737

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ccgaggaact ggccgagtag taggtagaga atcggttttt cttgtctcag cccggggtct 120

gctgcgtctg gtgggtggtga aggggagaaa ttcgtgagat ctgttccgga tcaggcgtgc 180

gagctcggga atcagggggtt tcacacatag cttcgtcgat ttgaatttga tgtactaatg 240

gagtctaagg gtggcaaaaa gtctagcagt agtcgttcta tgatgtatga agctcccttg 300

gctacagcat tgaggacgtt cgacctgccg gaggcntnaa gaagttccag tctgctgctt 360

actncaactg cgcgaaaaan ccattctgat atcctttggc tttcctcatc taatagttta 420

ngattctttt ctgacctttg at 442

<210> 738

<211> 527

<212> nucleic acid

<213> Zea mays

<400> 738

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acccacgcgt ccgaaagaat tccctcccct agccgccgcc gccgctcgtc tccctgagag 120

gtgccgggac cgaggatttg gncgagttct aggtagagaa tcggttttac ttgtctcagc 180

ccgggggtctg ctgcgtctgg tgggtggtgaa ggggagaaat tcgtgagatc tgttccggat 240

caggcgtgcg agctcgggaa tcaggggttt cacacatagc ttcgtcgatt tgaatttgat 300

gtactaatgg agtctaaggg tggcaaaaag tctagcagta gtcgttctat gatgtatgaa 360

gctncccttg gctacagcat tgaagacgtt cgacctgccn gaagcctgaa aaagttcaat 420

ctgntgctta ctccaactgc gccaaagaaa ccattctgat atcccttttg gcttccctcat 480

tctaagtaag tttaaggatt cttttctgac actttgattc ttancaa 527

<210> 739

<211> 249

<212> nucleic acid

<213> Zea mays

<400> 739

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ccaagctggg acatggggaa aggaacttgg tgcggaggct tatgactgca acaacatggt 120  
cgagcaggag ctgcctggag gtgggatcct catctaccag agcttctgtg ctgctgaaga 180  
cgccgttgct agctcgccca aatccgttct tcgctgcttt gatggcgaga atgcagcact 240  
tttgcaag 249

<210> 740  
<211> 514  
<212> nucleic acid  
<213> Zea mays  
<400> 740

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accacgcgt ccggttgcca aagaattccc tcccctagcc gcgcgcgcgc ctgctctccc 120  
tgagaggtgc cgggaccgag gatttgccg agttctaggt agagaatcgg ttttacttgt 180  
ctcagcccg ggtctgctgc gtctggtggt ggtgaagggg agaaattcgt gagatctgtt 240  
ccggatcagg cgtgcgagct cgggaatcag gggtttcaca catagcttcg tcgatttgaa 300  
tttgatgtac taatggagtc taagggtggc aaaaagtcta gcagtagtcg ttctatgatg 360  
tatgaagctc cccttggtcta cagcattgag gacgttcgac ctgccggagg cgtgaagaag 420  
ttccagtctg ctgcttactc caactgcgcg aagaaccatc ctgatatccc ttttggcttn 480  
ctcattctag taanttaaga attcntttnt tgac 514

<210> 741  
<211> 298  
<212> nucleic acid  
<213> Zea mays  
<400> 741

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tcctgacaat cgaccagttc ccagtccttg ctccctgcag tctccctcc tccatctcca 120  
gcgttggtgt ctgactcacc tgcaccaatg gctgttcttt ctgctgctgg tgctcccccg 180  
gcctcagcta tcgggtttga gggctatgag aagcgcttg agatcacatt ctctgaggca 240  
cctgtctttg tggaccccat ggcagcggtt tgcgtgccct ctccaggctc cagattga 298

<210> 742



<211> 217  
 <212> nucleic acid  
 <213> Zea mays

<400> 742

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 tctccatagc ttcatgctca ttgtggtaga cgcagctgac cacctagagc tgcatactta 120  
 cttattcaga ctgatccagc atgtggcaac atcggaagtt cacatgctta cagacgacaa 180  
 attttcataa caatagcctg gcttggctgc ttcacag 217

<210> 743  
 <211> 181  
 <212> nucleic acid  
 <213> Zea mays

<400> 743

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 tccacaatcc atgtgacgcc cgaggacggt ttcagctacg ccagttacga gggttatgggc 120  
 ttggatgcca ctgctctgtc ttatggtgac cttgtcaaga gggtcctccg gtgctttggc 180  
 c 181

<210> 744  
 <211> 373  
 <212> nucleic acid  
 <213> Zea mays

<400> 744

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 ctgctctctg tggtggtgaa ggggagaaat tcgtgagatc tgttccggat caggcgtgctg 180  
 agctcgggaa tcaggggttt cacacatagc ttcgtcgatt tgaatttgat gtactaatgg 240  
 agtctaaggg tggcaaaaag tctagcagta gtcgttctat gatgtatgaa gctccccttg 300  
 gctacagcat tgaggacgtt cgacctgccg gaggcgtgaa gaagttccag tctgctgctt 360  
 actccaactg cgc 373

<210> 745

<211> 281  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 745  
  
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 tgaatttnat gtactaatgg agtctaaggg tngcaaaaag tctagcagta gtcgtttctat 180  
 gatgtatgaa gctcccccttg gctacagcat tgaggacgtt cgacctgccg gagngtgaa 240  
 gaagttccag tctgctgctt actccaactg cgcgaagang c 281

<210> 746  
 <211> 184  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 746  
  
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 ttgaggacgt tcgaactgcc ggaggggtgaa gaagttccag tctgctgctt actccaactg 120  
 cgcgaagaag ccatactgat atcccttttg gcttccctcat tctagtagtt taggatttct 180  
 tttc 184

<210> 747  
 <211> 170  
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 <400> 747  
  
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 tggttaacct tgagatgtgc atgactggtc tggacaagaa gaaagcttgt gtctttttca 120  
 agactaatgc tgatgggaac acaacatgtg ccaaggaaat gacaaagctc 170

<210> 748  
 <211> 441  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 748

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 ggtggtggtg aangggagaa attcntgaga tctgtncogg atcaggcgng cgagctcggg 180  
 aatcaggggt ttcacacata gcttcgtccg atttgaattt gatgtactan atggantcta 240  
 anggtggcaa aaagtctanc attagtcgct ctatgatgta tgaanctccc ctnggctaca 300  
 gcattgagga cgttcgacct gccggaggcg tgnanangtt ccagtctgat gcttactnca 360  
 actgcgcgan ganccatcct gatatacctt ttggntncct cattctagna cgtaggant 420  
 tnttggtgac acttngattc t 441

<210> 749  
 <211> 195  
 <212> nucleic acid  
 <213> Zea mays

<400> 749

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 gactaangct gatgggnaca cagcatgtgc naaggaatga caaagctctc tggcatctct 180  
 gagtcatccc cgagt 195

<210> 750  
 <211> 463  
 <212> nucleic acid  
 <213> Zea mays

<400> 750

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 gccgccgccg ctgctctccc tgagaggtgc cgggaccgag gatttggccg agttctaggt 120  
 agagaatcgg ttttacttgt ctacgcccg ggtctgctgc gtctggtggt ggtgaagggg 180  
 agaaattcgt gagatctgtt ccggatcagg cgtgcgagct cgggaatcag gggtttcaca 240  
 catagcttcg tcgatttgaa tttgatgtac taatggagtc taagggtggc aaaaagtcta 300  
 gcagtagtcg ttctatgatg tatgaagctc cccttggtta cagcattgag gacgttcgac 360  
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tgataatcctt ttnggtttct taatctaaga nnttangatt ctt

463

<210> 751  
<211> 292  
<212> nucleic acid  
<213> Zea mays

<400> 751

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cttgtggacc tcatgggcgt ggtttgctg nccctctnccn agggcccaga ttgaactctg 180

tacnggatct tggcaggng caaaattgtg tctggagctc tccaacaagg atttcgactc 240

atatgtcctn tctgaagtna agcttgnta tcnnnccag gaagatngac ca 292

CCAGGTTTCA  
CACATAATNT  
AGCAGTAGTC  
ACCTGCCGGA  
TCCTGATATC  
CCTTTTGGCT  
TCCTCGTTCT  
AGTAGTTTAG  
GATTTCTTTT  
C

<210> 752  
<211> 291  
<212> nucleic acid  
<213> Zea mays

<400> 752

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cacataatnt nactanttga atttgatgta ctaatggant ctaanggtgg caagaagtnc 120

agcagtagtc gntccatgat gtatgaagct ccccttggt acagcatnng aanncgttcg 180

acctgccgga gnncaagaa gttccagtct gctgcttact ccaactgcgc aaagaagcca 240

tcctgatatc ccttttggct tcctcgttct agtagtttag gatttctttt c 291

<210> 753  
<211> 506  
<212> nucleic acid  
<213> Zea mays

<400> 753

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gaagaattcc ctcatagccg ccgccgccgc cgccgccgcc gctcatctcg ccgagaggtg 120

acgggaccga ggaactcgcc gagttctaag atagagaatc gcttttcaca gcctggggtc 180

tgctgcgtag ggtggtgttg aaggggagaa gtttaagatc tgttccacag atcacgcgtg 240

cgcgctcgcg aatcgggggt tccacacata gcttcgctcg tttgaatttg atgtactaat 300  
ggagtctaag ggtggcaaga agtctagcag tagtcgttcc atgatgtatg aagctcccct 360  
tggctacagc attgaggacg ttcgacctgc cggaggcgcc aagaagttcc aatctgctgc 420  
ttaactccaac tggcgcaaag aaancatcct gaaaaccctt ttggnntcct ccttccaata 480  
atttaaggat tccttttcna aaacnt 506

<210> 754  
<211> 275  
<212> nucleic acid  
<213> Zea mays

<400> 754

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cccttttggc ttctcattc tagnagtta ggatttcttt totgacactt tgatgctgac 180  
naatctctct ggctgggtgc ttctgataa tcgaccagtt cccagtcttg ctcttgcac 240  
cctcgtctct ccattctncag cattgtgtcc gattc 275

<210> 755  
<211> 158  
<212> nucleic acid  
<213> Zea mays

<400> 755

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gctctaggtg gtcagctgcg tctaccacaa tgagcatg 158

<210> 756  
<211> 165  
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<213> Zea mays

<400> 756

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tgctgatgct tccccggtct cagctatcgg gtttgagggc tatgagaagc gccttgagat 120

cacattctct gaggcacctg tctttgtnga ccttcattcn cntng 165

<210> 757  
 <211> 157  
 <212> nucleic acid  
 <213> Zea mays

<400> 757

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 tgcgtgccct ctccagggcc cagattgact ctgttctgga tcttgcaagg tgcacaattg 120  
 tgtccgagct ctccaacaag gatttcgact catatgt 157

<210> 758  
 <211> 221  
 <212> nucleic acid  
 <213> Zea mays

<400> 758

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 totggcactc tgaaatcatc cccgagatgg agatctgcga ttttgacttc gaaccctgtg 120  
 gctaactccat gaatgcgata catggntctg cattctccac aanccatgtg acgcccagg 180  
 acggtttnag ctangcnagt tangaggtn tgggctgggn n 221

<210> 759  
 <211> 493  
 <212> nucleic acid  
 <213> Zea mays

<400> 759

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 cgggaccgag gaactggccg agtactaggt agagaatcgg ttttcttgt ctcagccggg 180  
 gtctgctgcg tctggtggtg gtgaagggga gaaattcgtg agatctgttc cggatcggcg 240  
 tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc gatttgaatt tgatgactaa 300  
 tggagtctaa gggtagcaaa aagtctagca gtagtcgttc taatgatgta tgaactcccc 360  
 ttggctacag cattgangac gttcgacctg ccggaggcgt naagaanttc cagcctgctg 420

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nttaaggatt tct 493

<210> 760  
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<212> nucleic acid  
<213> Zea mays  
  
<400> 760

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<213> Zea mays  
  
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tgcaacaaca atggtggaat aggancttcc tggaaggcgg ggntcctcgt gtaacaaaaa 180  
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<210> 764

<211> 147

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<213> Zea mays

<400> 764

agcttgatga gtaagacggg cttctggtgt cgatttgctt ctgagttgtt tattttatat 60  
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<210> 765

<211> 187

<212> nucleic acid

<213> Zea mays

<400> 765

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gaggtctatg actgcaacaa catgtggagc aggagctgcc tggaggcggg ctctcgtat 120  
accagagctt ctgcgctgct gaagacgtg ttgctacctt cgcccaaate tgttttccac 180  
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<210> 766

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<212> nucleic acid

<213> Zea mays

<400> 766

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ccagcattgt gttctgattc acctgctcca atggctgttc tttctgctgc tgatgcttcc 120  
ccggtctcag ctatcgggtt tgaggcatg agaa 154



<210> 767  
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 ggcactacca agctcctgct caccattcca agaatccttg ag 162

<210> 768  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 ggtggtgaag gggagaaaatt cgtgagatct gttccggatc aggcgtgcga gctcgggaat 180  
 caggggtttc acacatagct tcgtcgattt gaatttgatg tactaatgga gtctaagggt 240  
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<210> 769  
 <211> 295  
 <212> nucleic acid  
 <213> Zea mays  
  
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 gagatctgtt ccggatcagg cgtgcgagct cgggaatcag gggtttcaca catagcttcg 180  
 tcgatttgaa tttgatgtac taatggagtc taagggtggc aaaaagtcta gcagtagtcg 240  
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<210> 770

<211> 329  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 770  
  
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 atgactgcaa caacatgggtg gagcagagct gcctggagcg ggctctctgt accagagttc 120  
 ngtgctgctg aagacgtgtg ntactcgcca atctgttncc actgcttgac ggcgagacgt 180  
 ggaagtgtcc tcctcctatg aagaaggata caagctggct aatcttctct gtgggaggag 240  
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 ttnttctgag tgttattnn cgcgcaatt 329

<210> 771  
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 <400> 771  
  
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 ggtcgaccca cgcgtccgat tnantcccat agccgacnac ggcgntcgtc tgctgagag 120  
 gtgccgggac cgaggnactg gncgagttct agggagagaa tcggttttgc ttgtctnagc 180  
 acggggtctg ntgcgtctgg tgggtggtgaa ggggagaaat tcgtgagatc tgttccggat 240  
 caggcgtgcg agctcgggaa tcaggggttt cacacatagn ttcgtngatt tgaatttgat 300  
 gtactaatgg agtctaaggg tggcaaaaag tctagcangg agtcgttcca tgatgtatga 360  
 agtccccctt ggntacagga ttgaagacgt tagactncct ganggcatga aaaaagttca 420  
 agctggnggt tnnttcaanc nccncnanaa accgttctn aattcccttt nggttnctta 480  
 atctnaaann ttaaganttn ctttcttaaa a 511

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gctctaggtg gtcagctgcg tctaccacaa tgagcatgta cgtatggaga aatatctgtg 120  
 aaoccatttt gtttatgaat aagatttggtt tttt 154

<210> 773  
 <211> 160  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 773

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 acccattttg tttatgaata agatttggtt ttttcgagtt 160

<210> 774  
 <211> 126  
 <212> nucleic acid  
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 <400> 774

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 cacatt 126

<210> 775  
 <211> 391  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 775

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 ccgggaccga ggaactggcc gagttctagg tagagaatcg gttttgcttg tctcagcacg 180  
 gggctctgctg cgtctgggtg tgggtgaagg gagaaattcg tgagatctgt tccggatcag 240  
 gogtgcgagc tcgggaatca ggggtttcac acatagcttc gtcgatttga atttgatgta 300  
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 ccccttggct acagcattga ggacgttcga c 391

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 ccgangaact ggccgngttc taggtagaga atnggttttg cttgtctcan cacgggggtct 180  
 gctgcgtctg gtgggtggtga aggggagaaa ttcgtgagat ctgttccgga tcaggcgtgc 240  
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<210> 779

<211> 117

<212> nucleic acid

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<210> 780

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gaatcagggg ttccacacat agctcgtcga tntgaatttg atgtactaat ggagtctaag 180

ggtggcaaaa agtctagcag tagtcgttct atgatgtatg aagctcccct tggctacagc 240

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agatctgttc cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt 180

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<211> 272

<212> nucleic acid

<213> Zea mays

<400> 783

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cgatttgaat ttgatgtact aatggagtct aagggtgcaa aaagtctagc agtagtcgtn 240

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agatctgttc cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt 180  
cgatttgaat ttgatgtact aatggagtct aagggtnngn aaaagtctag cagtagtcgt 240  
ctatgatgta tna 253

<210> 823  
<211> 269  
<212> nucleic acid  
<213> Zea mays

<400> 823

caaagaattc cctcccctag ccgccgccgc cgctcgtctc cctgagaggt gccgggaccg 60  
aggaactggc cgagttctag gtagagaatc ggttttgctt gtctcagcac ggggtctgct 120  
gcgtctggtg gtggtgaagg ggagaaattc gtgagatctg ttccggatca gccgtgcgag 180  
ctcgggaatc aggggtttca cacatagctt cgtcgatttg aatttgatgt actaatggag 240  
tctaagggtg gcaaaaagtc tagcagtag 269

<210> 824  
<211> 296  
<212> nucleic acid  
<213> Zea mays

<400> 824

cttcttgttg ccaaagaatt cctccccta gccgccgccg ccgcttcgtc tccctgagag 60  
gtgccgggac cgangaactg gccgagttct aggtagagaa tcggttttgc ttgtctcagc 120  
acgggggtctg ctgcgtctgg tgggtgtgaa ggggagaaat tcgtgagatc tgttccggat 180  
caggcgtgag agctcgggaa tcaggggttt cacacatagc ttcgtcgatt tgaatttgat 240  
gtactaatgg agtctaaggg tggcaaaaag tctagcagta gtcttccatg atgtat 296

<210> 825  
<211> 296  
<212> nucleic acid  
<213> Zea mays

<400> 825

gncaaagaat tccctcccct agccgccgcc gccgctcgtn tccctgagag gtgccgggac 60  
cgaggaactg gccgagttct aggtagagaa tcggtnttgc ttgtctcagc acgggggtctg 120

ctgcgtctgg tgggtggtgaa ggggaganat tcgtgagatc tgttcoggat caggcgtgcg 180  
 agcncgggaa tcagggggttt cacacatagc tcgtcgattt gaatttgatg tactaatgga 240  
 gtctaagggt ggcaaaaagt ctagcatagt cgttccatga tgtatgaagc tcccct 296

<210> 826  
 <211> 267  
 <212> nucleic acid  
 <213> Zea mays

<400> 826

aaaagaattc cctcccctag ccgccgccgc cgctcgtctc cctgagaggt gccgggaccg 60  
 aggaactggc cgagttctag gtagagaatc ggttttgctt gtctcagcac ggggtctgct 120  
 ggtctggtgg tgggtgaaggg gagaaattcg tgagatctgt tccggatcag gcgtgcgagc 180  
 tcgggaatca ggggtttcac acatagcttc gtcgatttga atttgatgta ctaatggagt 240  
 ctaagggtgg caaaaagtct agcagta 267

<210> 827  
 <211> 506  
 <212> nucleic acid  
 <213> Zea mays

<400> 827

gagcggnnnn tnnntatggt ggggggnnta atnnttctac tgggtccggaa ttcccgggtc 60  
 gacccacgcg tccggttgcc aaagaattcc ctcccctagc cgccgccgcc gctcgtctcc 120  
 ctgagaggtg ccgggaccga ggatttgccc gagttctagg tagagaatcg gttttacttg 180  
 tctcagcccg gggctctgctg cgtctggtgg tgggtgaaggg gagaaattcg tgagatctgt 240  
 tccggatcag gcgtgcgagc tcgggaatca ggggtttcac acatagcttc gtcgatttga 300  
 atttgatgta ctaatggaat ctaanggtgg caaaaantct aacaataatc gntctatgaa 360  
 tggatgaaac tncccttggc tncancattg aagaccttcg anctnnccga agcnttaaaa 420  
 aattccagtc tgntggttac ttcaactnng ccgaaaaacc attctggaat tccctttnng 480  
 tttcctnaat ctaatannta aggatt 506

<210> 828  
 <211> 295  
 <212> nucleic acid

<213> Zea mays

<400> 828

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gagaggtgcc gggaccgagg aactggccga gttctaggta gagaatcggg ttgcttgctc 120  
tcagcacggg gtctgctgcg tctggtggtg gtgaagggga gaaattcgtg agatctgttc 180  
cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt cgatttgaat 240  
ttgatgtact aatggagtct aagggtggca aaaatctagc agtagtcgtt ccatg 295

<210> 829

<211> 265

<212> nucleic acid

<213> Zea mays

<400> 829

gttgccaaag aattccctcc cctagccgcc gccgccgctc gtctccctga gaggtgccgg 60  
gaccgaggaa ctggccgagt tctaggtaga gaatcggttt tgcttgcttc agcacggggg 120  
ctgctgcgtc tgggtggtgg gaaggggaga aattcgtgag atctgttccg gatcaggcgt 180  
gcgagctcgg gaatcagggg ttccacacat agcttcgtcg atttgaattt gatgtactaa 240  
tggagtctaa ggggtggcaaa aagtc 265

<210> 830

<211> 271

<212> nucleic acid

<213> Zea mays

<400> 830

cttcttggtg ccaaagaatt ccctccccta gccgccgccg ccgctcgtct ccctgagagg 60  
tgccgggacc gaggaactgg ccgagttcta ggtagagaat cggttttgct tgtctcagca 120  
cggggctctgc tgcgtctggt ggtggtgaag gggagaaatt cgtgagatct gttccggatc 180  
aggcgtgcga gctcgggaat caggggtttc acacatagct tcgtcgattt gaatttgatg 240  
tactaatgga gtctaagggt ggcaaaaagt c 271

<210> 831

<211> 228

<212> nucleic acid

<213> Zea mays

<400> 831

tcattctcgcc gagaggtgac gggaccgagg aactcgccga gttctaagat agagaatcgc 60  
ttttcacagc ctgggggtctg ctgcgtaggg tgggtgttgaa ggggagaagt ttaagatctg 120  
ttccacagat cagcgtgcg cgctcgcgaa tcgggggttc cacacatagc ttcgtcgttt 180  
tgaatttgat gtactaatgg agtctaaggg tggcaagaag tctagcag 228

<210> 832

<211> 227

<212> nucleic acid

<213> Zea mays

<400> 832

tcgtctccct gagaggtgcc gggaccgagg aatggccgag tactaggtag agaatcggtt 60  
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gatctgttcc ggatcaggcg tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc 180  
gatttgaatt tgatgtacta atggagtcta aggttgcaa aaagtct 227

<210> 833

<211> 270

<212> nucleic acid

<213> Zea mays

<400> 833

gttgccaaag anttccctcc cctagccgcc gccgccgctc gtactccctg agaggtgccg 60  
ggaccgagga actggccgag ttctaggtng agnatcggtt ttgcttgtct cagcacgggg 120  
tctgctgctg ctggtggtgg tganggggag aaattcgtga gatctgttcc ggatcaggcg 180  
tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc gatttgaatt tgatgtacta 240  
atggagtcta anggtggcaa aaagtctanc 270

<210> 834

<211> 222

<212> nucleic acid

<213> Zea mays

<400> 834

tcgtctccct gagaggtgcc gggaccgagg aactggccga gtactaggta gagaatcggg 60  
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 agatctgttc cggatcaggc gtgcgagctc gggaatcagg ggtttcacac atagcttcgt 180  
 cgatttgaat ttgatgtact aatggagtct aagggtggca aa 222

<210> 835  
 <211> 261  
 <212> nucleic acid  
 <213> Zea mays

<400> 835

aaagaattcc ctcccctagc cgcgcngcc gctcgtctcc ctgagaggtg ccgggaccga 60  
 ggaactggcc gagttctagg tagagaatcg gttttgcttg tctcagcacg gggctctgctg 120  
 cgtctgggtg tggtgaagg gagaaattcg tgagatctgt tccggatcag gcgtgcgagc 180  
 tcgggaatca ggggtttcac acatagcttc gtcgatttga atttgatgta ctaatggagt 240  
 ctaagggtgg caaaaatcta c 261

<210> 836  
 <211> 97  
 <212> nucleic acid  
 <213> Zea mays

<400> 836

tcagctgcgt ctaccacaat gagcatgaac gtatgngac atatctgtga acccattttg 60  
 tttatgaata agatttggtt ttntcgagtn aaaaaaa 97

<210> 837  
 <211> 229  
 <212> nucleic acid  
 <213> Zea mays

<400> 837

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 tgcttgtctc agcacggggt ctgctgcgtc tgggtgggtg gaaggggaga aattcgtgag 120  
 atctgttccg gatcaggcgt gcgagctcgg gaatcagggg tttcacacat agctcgtcga 180  
 tttgaatttg atgtactaat ggagtctaag ggtggcaaaa agtctagca 229

<210> 838  
 <211> 254  
 <212> nucleic acid  
 <213> Zea mays

<400> 838

ganttcctc ccctagccgc cgccgccgct acgtctccct gagagggtgcc gggancgagg 60  
 aactggccga gttctaggta gagaatcggg tttgcttgct tcagcacggg gtctgctgcg 120  
 tctgggtggtg gtgaagggga gaaattcgtg agatctgttc cggatcaggc gtgcgagctc 180  
 gggaatcagg ggtttcacac atagcttcgt cgatttgaat ttnatgtact aatggagtct 240  
 aagggtggca aana 254

<210> 839  
 <211> 493  
 <212> nucleic acid  
 <213> Zea mays

<400> 839

caaagaattc cctcccctag ccgccgncgn cgctcgtctc cctgaagaag gtgcccggga 60  
 ccgagggatt tggcccgagt tctaggtaag aagaaatcgg gttttacttg tctcaagccc 120  
 ggggggtctgc ttgcgtctgg tgggtgggtga aaggggaaga aattcgtgaa gatctgttcc 180  
 ggatcaggcg tgcgaanctc gggaatcagg gggtttcaca cataagcttc gtcgatttga 240  
 atttgatgta ctaaattgga agtctaaagg gtggcaaaaa gtctagcagt aagtccgttc 300  
 tatgatgtat gaaacttccc tttggctaca gcatttgaag gacgttcgac ctggccggan 360  
 gccttgaaga angttccagt ctgnttgntt acttcaactt gngccaaaaa accattctgg 420  
 atattccttt tgggnnttcc tcaatctagt annttaanga nttcttttct gacacttttg 480  
 aatcttganc caa 493

<210> 840  
 <211> 296  
 <212> nucleic acid  
 <213> Zea mays

<400> 840

angnacgcgt acgttagctc gnaattcngc tcgagcggct cgagcnaaga attccntccc 60



ctagccgccg ccgncgccgc tcgtctccct gagaggtgcc ggnaccgagg aactggccga 120  
 gtactaggta gagaatcgnt tttcttctgc tcagcccggg gtctgctgcg totggtggtg 180  
 gtgaanggga gnaattcgtn agatctgttc cggatcaggc gtgcgagctc gggaatcagg 240  
 ggtttcacac atagcttcgt cgatttgaat ttgatgtact aatggagtct aagggg 296

<210> 841  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 841

tngccnnana nttccnnccc ctagccgccn ccgcccgcgc tngtntccct gagaggtanc 60  
 cgggaccgat gaactngccg agtactaggt agagaatcgg tttntcttnt ctacagcccgg 120  
 ggtctgctnc ntctggtggt ggtgaagggg agaaattcgt ganatctgtt ccggancatg 180  
 cgtgngagct cnggaatcag gggtttcaca catagcttcg tcgatttgaa tntgatgtat 240  
 aatggatcta agggtnngcaa naantctagc agtagtcgtn cnagntgnat ga 292

<210> 842  
 <211> 254  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 842

cgccgccgct catctcgccg agaggtgacg ggaccgagga actcgccgag ttctaagata 60  
 gagaatcgct tttcacagcc tggggtctgc tgcgtagggg ggtgttgaag gggagaagtt 120  
 taagatctgt tccacagatc acgcgtgcgc gctcgcgaaat cgggggttcc acacatagct 180  
 tcgtcgtttt gaattgatgt actaatggag tctaaagggtg gcaagaagtc tagcatatcg 240  
 ttccatgatg tatg 254

<210> 843  
 <211> 255  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 843

gccaaagaat tccctccctt agccgccgcc gccgctcgtc tccctgagag gtgccgggac 60

cgaggaactg gccgagttct aggtagagaa tcggttttgc ttgtctcagc acggggtctg 120  
 ctgcgtctgg tgggtggtgaa ggggagaaat tcgtgagatc tgttccggat caggcgtgcg 180  
 agctcgggaa tcaggggttt cacacatagc ttcgctgatt tgaattgatg tactaatgga 240  
 gtctaagggt ggcaa 255

<210> 844  
 <211> 214  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 844

tcctctcgcc gagagggtgac gggaccgagg aactcgccga gttctaagat agagaatcgc 60  
 ttttcacagc ctgggggtctg ctgcgtaggg tgggtgttgaa ggggagaagt ttaagatctg 120  
 ttccacagat cagcgtgcg cgctnccga atcggggggtt ccacacatag cttcgtcgtt 180  
 tgaatttgat gtactaatgg agtctaaggg tggc 214

<210> 845  
 <211> 798  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 845

gccacgcgtg cgncaaaaca acaatgnntn tnntttggag ggcttttgaa aaagcccctt 60  
 gggggatcaa cattcttctt gagggcnccc tggcttttg tgggaccct catgggcgtg 120  
 ggttttgncg tgccctctt ccaaggcccc agattgactc tgttctggga tctttgcacg 180  
 ggngcacaaa ttgtgtcccg agctcttcca acaagggtt tccgactcat atgtcctttc 240  
 tgaggcaaag cttgnntatc tctctctga agattgtcat caaagacctg tggcactacc 300  
 aagctcctgc tcaccattcc aaagaatcct tgaacttgct tgaaaaactt gtctatgcc 360  
 ctttgctgct tgtgaaaata ctcccgtagg acggtcatct tttctggng cacagncagg 420  
 cccccacaa ggaacttctt ttgaaggaag ttgntgnact ttaaaccgct tcttttggcc 480  
 ggcttgaaa tctggggggg naatgcntn tgtggaattg gggaatncca ncnaggaacc 540  
 tggacaanaa atnggnaccg ntttttaacn ccccttgatt ncccnagca acccaatggg 600  
 tnaacctna aaaggccnt gactgggctg ggcnnnnnnn nnnnnnnnnn nnnnnnnnaa 660

naataatgtt gangggaaca caacttgggc caagggaaat ganaaacttt ttgnnttttg 720  
 aaaatatncc ccgaanggaa aacngnantt tantttaacc cgggggttttc aggaaggggc 780  
 ccagggtttg ttttcnaa 798

<210> 846  
 <211> 268  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 846

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 ttttcttgtc tcagcccngg gtctgtctgog tctgggtggg gtgaanggga gaaattcgtg 120  
 aganctgtac cactcaggcg tgcgagctcg ggaatcaggg gtttcacaca tagcttcgtc 180  
 gatttgaatt tgatgtacna tggagtctaa ggggtggcaaa aatctagnen tatcgttcta 240  
 ngattatnaa gccccctngg taaannat 268

<210> 847  
 <211> 276  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 847

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 ggtagagaat cggtttttct tgtctcagcc cgggggtctgc tgcgtctggg ggtgggtgaag 120  
 gggagaaatt cgtgagatct gttccggatc angcgtgcga gtcgaggat caggggtttc 180  
 acacatagct cgtcgatttg aatttgatgt acnaatggat cnaaggggtg caaaaagtct 240  
 agcagnagtc gtctatgatg tatgaagctc cccttg 276

<210> 848  
 <211> 185  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 848

gcaagctacc tgtgacaaaa acgattccaa gaaaccactt cttgtgcttc ctccactggc 60  
 tgtctgtgca tcttgctcta aaacaatcgc agtcctgcaa gttgttgctg ttgctcctac 120

ccctgcctct gcaattgggt ttgagggata tgagaagcgc ctcgagatca gcttctatga 180  
ggcac 185

<210> 849  
<211> 519  
<212> nucleic acid  
<213> Zea mays  
  
<400> 849

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ccgggtcgac ccacgcgtcc gagagcttcg actcatatgt tctgtccgag tccagcctct 120  
ttgtgtatcc gggacagggg tgtttctgaag acctgtggga cgacgaagct gctgctcgcc 180  
attcctcgca tccttgagct tgetgcaggg ctgtcactgc ctcttctttc agcgaagtac 240  
tctcgtggga cttcatctt ccccggcgcg cagcccgcg caccgcgag cttctcggag 300  
gaggtatctg tgctgaacgc tttctttggg aacctcaagt ccggtggcaa tgcctacctg 360  
atcgggtgaac cgttcaagcc caacaagaag tggcatgtct actacgccac agaggagcct 420  
gagcgtccta tggtgacgct tgagatgtgc atgacagagc ttgacgttaa gaaagctgct 480  
gtgttcttca agaacttcac tgggtggcaac ttnacntna 519

<210> 850  
<211> 324  
<212> nucleic acid  
<213> Zea mays  
  
<400> 850

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gtggcctgcg cgccctctcc cgtgagcaga tcgactcgtt tctggatctc gcacggtgca 120  
ccatagtgtc acagctctcc aacgagagct tcgactcata tgttctgtcc gagtccagcc 180  
tctttgtgta tccccacaag gttgttctga agacctgtgg gacgacgaag ctgctgctcg 240  
ccattcctcg catccttgag cttgctgcag ggctgtcact gcctcttctt tcagcgaagt 300  
actctcgtgg gaccttcac ttcc 324

<210> 851  
<211> 260  
<212> nucleic acid

<213> Zea mays

<400> 851

gaagcgccctc gagatcacgt tctctgacgc gcctgtcttt gaggaccctt gtggtcgtgg 60  
cctgcgcgcc ctctcccgtg agcagatcga ctcgtttctg gatctcgac ggtgcacccat 120  
agtgtcacag ctctccaacg agagcttcga tcatatgtt ctgtccgagt ccagcctctt 180  
tgtgtatccc cacaaggttg ttctgaagac ctgtgggacg acgaagctgc tgctcgccat 240  
tcctcgcatc cttgagcttg 260

<210> 852

<211> 272

<212> nucleic acid

<213> Zea mays

<400> 852

agcgccctcga gatcacgttc tctgacgcgc ctgtctttga ggacccttgt ggtcgtggcc 60  
tgcgcgccct actcccgtga gcagatcgac tcgtttctgg atctcgacg gtgcaccata 120  
gtgtcacagc tctccaacga gagcttcgac tcatatgttc tgtccgagtc cagcctcttt 180  
gtgtatccca caaggttggt ctgaagacct gtgggacgac gaagctgctg ctcgccattc 240  
ctcgcatcct tgagcttgct gcaggctgtc at 272

<210> 853

<211> 239

<212> nucleic acid

<213> Zea mays

<400> 853

gcgcctcgag atcacgttct ctgacgcgcc tgtctttgag gacccttgtg gtcgtggcct 60  
gcgcgccttc tcccgtgagc agatcgactc gtttctggat ctgcgacggg gcaccatagt 120  
gtcacagctc tccaacgaga gcttcgactc atatgttctg tccgagtcca gcctctttgt 180  
gtatccccac aaggttggtc tgaagacctg tgggacgacg aagctgctgc tcgccattc 239

<210> 854

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 854

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agtgtctcag ctctnccaan gagagctnag actcatatgt tctgtncgag tccagcctct 120  
ttgtgtatcc ccacaaggnt gttctgaaga cctgtggnac gacgaagctg ctgctcgca 180  
ttcctcgcat ccttgagctt gctgcagggc tgtcantgcc tcttctttca gcgaagtact 240  
ctcgtgggac cttcatcttc cccggcg 267

<210> 855

<211> 295

<212> nucleic acid

<213> Zea mays

<400> 855

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gaggaccctt gtggtcgtgg cctgcgcgcc ctctcccgtg agcagatcga ctggtttctg 120  
gatctcgcac ggtgcaccat agtgtcacag ctctccaacg agagcttcga ctcatatgtt 180  
cagtccgagt ccagcctctt gtgtatcccc acaaggttgt tncgaagact gtgggacgac 240  
gaagctgctg ctgccattc ctgcacatnt gagntgctgc aggctgtcat gcctc 295

<210> 856

<211> 266

<212> nucleic acid

<213> Zea mays

<400> 856

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cctctttgtg tatccccaca aggttggttct caagacctgt gggacnacga agctgctgct 120  
cgccattcct cgcaccttg agcttgctgc agggctgtca ctgcctcttc tttcagcgaa 180  
gtactctent gggaccttca tcttccccgg cgcgcagccg cgccacaccg cagcttctcg 240  
gaggaggtat ctgtgctgaa cgcttt 266

<210> 857

<211> 316

<212> nucleic acid

<213> Zea mays

<400> 857

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gctcgccatt cctcgcatcc ttgagcttgc tgcagggctg tcaactgcctc ttctttcagc 120  
gaagtactct cgtgggacct tcatcttccc cggcgcgcac cgcgcgcaca ccgcagcttc 180  
tcggaggagg tatctgtgct gaacgctttc tttggtaacc tcaagtccgg tggcaatgcc 240  
tacctgatcg gtgaaccgtt caagcccaac aagaagtggc atgtctacta cgccacagag 300  
gagcctgagc gtccta 316

<210> 858

<211> 219

<212> nucleic acid

<213> Zea mays

<400> 858

ggtatcttaa acttagaact cttcagcgtg catcatatac tcattacttc tatgaagcaa 60  
ctgttggtgc tggcctccct atcattagca ctctgcgtgg tctcctagag acaggtgaca 120  
agatattgcy cattgaaggg attttcagtga gaacgttaag ttatatatttt aacaacttcg 180  
aaggagcaag aaccttttagt gatgttggtg ctgaagcaa 219

<210> 859

<211> 323

<212> nucleic acid

<213> Zea mays

<400> 859

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tcgagcaata gctcaagggt gcagtgaata caatattact gttgtattga agcaacaaga 120  
ttgtgtaagg gctctgagag ctgctcactc aaggttcttc ctctccaaaa ccacacttgc 180  
tgttgggatac attggacctg gtttaattgg tggagcactc cttaaccagt tgaagaatca 240  
gactgctggt ctcaaggaaa atatgaacat tgatctgcgt gtcattggaa taactggctc 300  
aagtacaatg cttttgagcg aca 323

<210> 860

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 860

ctttctgata ttcctgtgaa gagccttgtg ccagaaacac ttgcagcatg ctcgtcagca 60  
gatgaattca tgcagaagct accatccttt gacgaggact gggccaggca acgcagttag 120  
tgctgatgcn gcagacgaag tcctgcgtta cgttggagtgt gtggatacgg tgaacaagag 180  
gggacagggtg gagctacgta ggtacaagag ggaccaccgg ttgcgcant ctccggctcc 240  
gacaacatca tcgcattcac cacctcgagg tacaaggagc agccgctaata cgtgcggngc 300  
ccaggcgcgg 310

<210> 861

<211> 242

<212> nucleic acid

<213> Zea mays

<400> 861

tgaacactct gtatgctttg ctgttccaga aaaggaagtt gcattggttt ctgcagcctt 60  
gcatgctagg ttctgtgaag cattggcagc agggagggtta tctaaggttg aagtcattcn 120  
taattgtagc atccttgcta ctgtcggcct gaggatggca agtacacctg gagtcagcgc 180  
gacccttttt gatgcactag ctaaggcaaa tnttaatgta agggcgatan tcaaggttgc 240  
ag 242

<210> 862

<211> 303

<212> nucleic acid

<213> Zea mays

<400> 862

tgaaaaacga cttgaaaaat ggttttctcg atgtccggct gagacaataa ttgctactgg 60  
gtttattgcg agtacacctg aaaacatccc aacaacctta aaaagagatg gaagtgattt 120  
ctctgcagcc ataattgggt ccttggttaa ggctcgccag gtgaccatct ggactgatgt 180  
tgatggggta tttagtgcag atcccagaaa agttagttag gctgtaatct taagcacatt 240  
gtcgatatcag gaggcctggg aaatgtcata ttttggggca aatgtcttgc atccacgtac 300  
tat 303



<210> 863  
 <211> 294  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 863  
  
 gtgaagtcag catatctttg aacttgatc catcgaaact atggagttgt gaattgggtc 60  
 agcacaaaaa tgaacttgat gatgtcattg aagagcttga gaagatagca gttgttcac 120  
 tactgcagaa caggtcaata atatcttta tttgaaatgt gcagagggtcg tctttgattc 180  
 ttgaaaaggc tttcaatgtt ctacgaagaa acggtgtcaa tgtccagatg atatcgcaag 240  
 gggcgtccaa ggtgaacatt tccttggtgg tccatgacag tgaggcaaaa cagt 294

<210> 864  
 <211> 224  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 864  
  
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 gaactgatac agcaggaact tgaccatgta gttgaagagc ttgagaaaat agcaattgtt 120  
 cgtctacttc agcagagggc gataatttca cttatcgga atgtggagca atcgtcnctc 180  
 atactagaaa agacgggacg tgtgctgagg aaaagtgggg ttat 224

<210> 865  
 <211> 240  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 865  
  
 tggagtgggtg gatgtggtca gcaagaaggg gcaggtggag ctgcgagcct acaagaggg 60  
 cccacccgtt cgcgcactat ctggttccga caacatcata gccttcacca cctcgaggta 120  
 caaggatcag cactgatcg tgcgagggtc gggggctggt gcggaagtga ctgcaggagg 180  
 cgtcttctgc gacatactgc gcctgtcttc ctaccttggt gcccctcgt agtctgtcaa 240

<210> 866  
 <211> 243  
 <212> nucleic acid

<213> Zea mays

<400> 866

cttgggatta agagaaattc aggtttggaa agatgttgat ggtgttttaa catgtgaccc 60  
 aaatatccat cctaaggcaa aacctgtgcc atacttgaca tttgatgagg cagctgaact 120  
 tgcctatfff ggagcgcagg ttttgcattc acaatcaatg cggccagcta gagaaggcga 180  
 tgtacctgtg agggttaaga atcttataac cgtcngccc ctggtaccct tatcataaag 240  
 cng 243

<210> 867

<211> 286

<212> nucleic acid

<213> Zea mays

<400> 867

gaaggtcata attcttgcaa gggaatctgg tttggggctt gaactttctg atattcctgt 60  
 acgaagtctt gtgccggang cttnaagtca tgtacctcag ccgacgaata catgcagaag 120  
 ttaccatctt ttgacgagga ttgggcaaga gagaggaaaa atgctgaagc tgcgggagaa 180  
 gttctgcggt atgttggagt ggtgnatgtg gtcncaaaaa gggccnaggt gaantnngcc 240  
 ctnaaagggg ggcccccent ttggnnaaaa ntggtcncaa aaaaaa 286

<210> 868

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 868

aagagaagct acaaactgaa gcagaacctg ccaatcttga taagtttggt catcatttgt 60  
 ctgagantca cttttttcct aacagagttt tggtagattg cacagcagat acaagtgttg 120  
 catctcacta ttatgattgg ctaaagaaag gaatccatgt catcactcca aacaagaaag 180  
 caaattcagg gccacttgac aggtatctta aacttagaac ttttcagcgt gcatcatata 240  
 ctcatctatt ctatgaagca actgttggtg ctggc 275

<210> 869

<211> 241

<212> nucleic acid



<400> 872

gtcattgatt cttgaaaagg cattcaacgt tctgcgaaga aatgggtgta atgtcnagat 60  
gatctcaciaa ggggcgtcca aggtgaacat atccttggtg gtcaatgaca gtgaggcaaa 120  
acagtgcgta caagccctcc attcagcgtt ctttgagaat ggattcttgt cggaggtaga 180  
gggagctgac gttccccaga acggagcctc tctgaactca aatggcgctg tctatggaaa 240  
ctagtcgact ctgcaactgc tttcttcaca atttgtgtac cattg 285

<210> 873

<211> 286

<212> nucleic acid

<213> Zea mays

<400> 873

ccaaggcgtc cgaaatccct gagctcgctg tcatcaagga cctgcatctt aggacggttg 60  
atgagctcgg tttagatagg tcgattgttt cagggtttatt ggatgaattg gagcaacttc 120  
ttaaaggagt tgctatgatg aaagagctga cccttaggac aagagattac cttgtttcct 180  
ttggtgaatg tatgtcgaca agaataattg ctgcatattt gaacaaaactt gggaaaaaag 240  
cacgccatat gatgcattga tatggcttat aaccaccgat gatttc 286

<210> 874

<211> 293

<212> nucleic acid

<213> Zea mays

<400> 874

ttcacaaatg cggatattct tgaagccact taccctgctg ttgcgaagag gctacatgaa 60  
gactggatgg atgaccctgc tattcctata gtcactggtt ttcttgggaa gggatgtaaa 120  
tcatgtgctg tcaccacttt aaggcagggg tggtagtgcac ttgacagcta caaccattgg 180  
caaagccttg gggattaaga gaaatccagg ttggaaggat gtagatggtg tgtgacatgt 240  
gatcctnata tatatggcaa atgctatanc ccgtgcctac ttgactttgg atg 293

<210> 875

<211> 266

<212> nucleic acid

<213> Zea mays

<400> 875

actgccaaagt cccttggcta tacagaacca gatccacgtg atgatctagg tggaatggat 60

gtcgctagaa aggcattgat cctggctagg cttcttgggc aacagataag catggagagt 120

atcaatgttg aggtttatat cctagtgaat taggacctga tgcagtgtcc acaaaggact 180

tcctgaaagt ggtttagtgc aattgacaag agcatggaag aaagaatcag agcagcatct 240

cgagaggaaa cgttctcgct acgtct 266

<210> 876

<211> 343

<212> nucleic acid

<213> Zea mays

<400> 876

ggctggttct aggacaaatg cttttaagtg atataggagt agatttgacc cagtggaaag 60

agaagctaca aactgaagca gaacctgcca atcttgataa gtttgttcat catttgtctg 120

agaatcactt tttccctaac agagttttgg tagattgcac agcagataca agtgttgcat 180

ctcactatta tgattggcta aagaaaggat ccatgtcatc actccaaaca agaagcaant 240

cagggccatt gacaggttct taancttgac tcttcagggg cataattatc atacttctan 300

gagnantgtg gnnngcccca nanatnnccn ngggtnnnan nag 343

<210> 877

<211> 442

<212> nucleic acid

<213> Zea mays

<400> 877

gngatctngn gnantancta ncttgngnan ntgngnggtn cacttnacnn cnannnncat 60

tggnanncn ttgagattaa ganatactcn tgttaggnaa gatgctgatg gtgntggaac 120

atgtgaccga gatatacatc cttaaagctga acctgancca tacttgacat ctngatgaat 180

cagctgaacn ngcctatddd ggagcgcagg aattgcatgc actatnnatg cggtcagngt 240

tgagaaggcg atntacctgt gagggctaata gaattnttat aactcgncta gnctanggta 300

cccttatcac taaaacntgn acgatatgag ttaagantgt tcagactngc attgtncctg 360

aaaatgaata ttnaatgctt gatatatcca gtacagggta tgctttgcca agtaccggat 420

naataanncc cntttaaggg aa

442

<210> 878  
<211> 285  
<212> nucleic acid  
<213> Zea mays

<400> 878

gnncaagacn tgtgccatac ttgacatttg atgaggcagc tgaacttgcc tatnttgag 60  
cgcagggtttt gcatccacaa tcaatgcggc cagctagaga aggcgatgta cctgtgaggg 120  
ttaagaattc ttataaccgt cganccctg gtacccttat cactaaagca agagatatga 180  
gtaagactgt ttgactagc attgttttga aatcaaatac tacaatgctt gatatatgca 240  
gcacacgcat gcttgccag tacggttttc tagctaaggt tttt 285

<210> 879  
<211> 282  
<212> nucleic acid  
<213> Zea mays

<400> 879

gncaaaacct gtgccatac ttgacatttg atgaggcagc tgaacttgcc tatnttgag 60  
cgcagggtttt gcatccacaa tcaatgcggc cagctagaga aggcgatgta cctgtgaggg 120  
ttaagaattc ttataaccgt cgagccctg gtacccttat cactaaagca agagatatga 180  
gtaagactgt ttgactagc attgttttga aatcaaatac tacaagctt atatagtcag 240  
cacacgcatg cttggccagt acggttttct agctaaggtt tt 282

<210> 880  
<211> 317  
<212> nucleic acid  
<213> Zea mays

<400> 880

gaaccagctg atattgatag tttgtttcat catttatccg ataatacagc atttccaaac 60  
aaagttttgg tggactgcac agcagataca agtggttgcac cccactatta taattgggta 120  
aagaagggtg ttcatgtcat cacaccaaatac aagaaagcaa attctggggc acttgatcag 180  
tacctgaaat tgcggactct gcaacgtgcc tcgtacactc attactttta tgaagcaact 240

gttggtgctg gcctcccaat cattagtacc ttgcgaggtc ttctggagac gggtgacaag 300  
 atactgcgta ttgaggg 317

<210> 881  
 <211> 315  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 881

ggatcaggtt gctgctctca aggaaaatat gaacattgat ctgcgtgtca ttggaataac 60  
 tggctcaagt acaatgcttt tgagcagaca cgggaataga cttaaccag tggaaacaac 120  
 ttctaaaaaa ggaagcagaa ccagctgata ttgatagttt tgttcacatc ttatccgata 180  
 atcacgtatt tccaaacaaa gttttggtgg actgcacagc agatacaagt gttgcatccc 240  
 actattataa ttggttaaag aaggggtattc atgtcatcac accaaataag aaagcaaatt 300  
 ctgggccact tgatc 315

<210> 882  
 <211> 288  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 882

aaccagctga tattgatagt tttgttcac atttatccga taatcacgta tttccaaaca 60  
 aagttttggt ggactgcaca gcagatacaa gtgttgcac ccactattat aattgggttaa 120  
 agaagggtat tcatgtcac acaccaaata agaaagcaaa ttctgggcca cttgatcagt 180  
 acctgaaatt gcgactctg caacgtgcct cgtacactca ttacttttat gaagcaactg 240  
 ttggtgctgg cctcccaatc atagtacctt gcgaggctct ctggagac 288

<210> 883  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 883

ccaaaacgac acttgacgtt ggcatcattg gacctggttt aattggtaga acactcctta 60  
 accagctcaa ggatcaggtt gctgctctca aggaaaatat gaacattgat ctgcgtgtca 120





cattatcctt gcaagggaaat caggtttgag gcttgaactt tctgatattc ctgtgaagag 180  
 ccttgtgcc aaaaacttg cagcatgctc gtcagcagat gaattcatgc agaagctacc 240  
 atcctttgac gaggactggg ccaggcaacg catgatgct 279

<210> 887  
 <211> 256  
 <212> nucleic acid  
 <213> Zea mays

<400> 887

tttantnntt tcgaaggaac aagggcattc agtngtgtcg ttgcngaagc aaaaganntg 60  
 ggatacactg aacctgaccc aaganntgat ctntctggaa ntgntgttgc ccgaaaagtc 120  
 attntccttg caaggggaatc aggnntgagg cttgaacttt ctggntattc ctgtgaagag 180  
 ccttgngetn gaaaacttg cagcagctcg tcagcagaga gttcagcaga agctaccatc 240  
 ctttgacgag gactgg 256

<210> 888  
 <211> 322  
 <212> nucleic acid  
 <213> Zea mays

<400> 888

gggagactgg atggatgacc ctgctattcc tatagtcact ggttttcntg gncagggatg 60  
 taaatcatgt gctgtcacca ctttaggcag gggtagtagt gacttgacag ctacaacccat 120  
 tggcaaagcc ttgggattaa gagaaatcca ggtttgaag gatgtagatg gtgtgttgac 180  
 atgtgatcct aatatatatg caaatgctat acccgtgcc tacttgactt ttgatgaggc 240  
 tgctgagctt gctactttg gtgcacaggt ttgcatccc caatcaatgc gaccggctag 300  
 ggatggtgat attccagtga ga 322

<210> 889  
 <211> 105  
 <212> nucleic acid  
 <213> Zea mays

<400> 889

gtgatcctaa tatatatgca aatgctatac ccgtgcccta cttgactttt gatgaggctg 60

ctgagcttgc ctactttggt gcacaggtta nacatcccc agtca 105

<210>	890
<211>	279
<212>	nucleic acid
<213>	Zea mays

<400> 890

gtgagtacaa cattactatt gtactgaagc aagaagactg cgtgcgggct ctgagagctg 60

ccactcaag gntcttctc tccaaaacca cactggctgn tggcatcatt ggacctgggn 120

tgattggtcg cacattactt aatcagctga aggatcaggc tgctgtcctc aaggaaaata 180

tgaacattga tttgcgtgtc atgggaatag ctggttctag gacaatgctt ttaagtgata 240

taggagtaga ttgacccag tggaaagaga agctacnaa 279

<210>	891
<211>	282
<212>	nucleic acid
<213>	Zea mays

<400> 891

gctatgcaaa tattaatgta agggcgatag ctcaaggttg cagtgaqtac aacattacta 60

ttgtactgaa gcaagaagac tgcgtgcggg ctctgagagc tgcccactca aggttcttcc 120

tctccaaaac cacactggct gttggcatca ttggacctgg gttgattggt cgcacactac 180

ttaatcagct gaaggatcag gctgctgtcc tcaaggaaaa tatgaacatt gtttgcgtgt 240

catgggaata gctggttcta ggacaatgct tttaagtgat at 282

<210>	892
<211>	282
<212>	nucleic acid
<213>	Zea mays

<400> 892

gttatccatg tttctgaaat cgaagagtgg aatatgggtca aaagcctaca tatcaagacg 60

gtggatgaac ttggacttcc aagatctgta atacaagaca tgctagatga actggagcaa 120

ctattgaaag gtatcgctat gatgaaagag ctgacgccta ggaccagtga ctaccttggt 180

tcatttggag aatgcatgtc caccaggatt ttttctgctt atttgaacaa aattcgtgtc 240

aaggcacggc agtatgacgc atttgatatt ggtttcatac aa 282

<210>	893
<211>	270
<212>	nucleic acid
<213>	Zea mays

<400> 893

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gtggatgaac	ttggacttcc	aagatctgta	atacaagaca	tgctagatga	actggagcaa	120
ctattgaaag	gtatcgctat	gatgaaagag	ctgacgccta	ggaccagtga	ctaccttggt	180
tcatttgag	aatgcatgtc	caccaggatt	tttctgctta	tttgaacaaa	attcgtgtca	240
aggcacggca	gtatgacgca	tttgatat	ttg			270

<210>	894
<211>	493
<212>	nucleic acid
<213>	Zea mays

<400> 894

gngnnnnnnng gnaantgttn anttngcccg ccagtagccga tccaagaatt cccgggtcga	60
cccacgcgtc cgatcagctt gggctagata gatcaattgt ttgtggtttg ttggatgaac	120
ttgaacaact tcttaaggga attgctatga tgaaagagct gactcttagg acacgagatt	180
atctagtttc atttggtgaa tgcattgtcta caagaatatt tgctgcactt ttaaacaaaa	240
ttggagtaaa agctcggcag tatgatgcat ttgatcttgg ctttataact accgatgatt	300
tcacaaatgc ggatattctg gaagcaactt atcctgctgt tgcaaggagg ctacatgtgg	360
aatggattaa tgatcctgct atacctatag tcaactggctt tcttggcaag ggatggagat	420
ctggtgcaat aactaccttg ggcanggggtg gtaatgactt gacagctaca accattggca	480
aaqccttggg aat	493

<210>	895
<211>	281
<212>	nucleic acid
<213>	Zea mays

<400> 895

aaaganctgn ctcttaggac acgagattat ctagttgcag tnggtgaatg catggctaca 60  
 agaanattgg cctgcacttt naaacaaaat tggagtaaaa gtctcggcag tatgatgcat 120  
 gtnatcttgg cttgataact accgatgatt tcacaaatgc ggntattctg gaagcaactt 180  
 atcctgctgt tgcaaggagg cgacatgtgg aatggattaa tgatcctgcn agacctatag 240  
 tcantggcnt tcttggcaag ggatggagat ctggtgcaat a 281

<210> 896  
 <211> 299  
 <212> nucleic acid  
 <213> Zea mays

<400> 896

gcgcggggcgc ggaggtgact gggggtggcg tcttctgcga catcctgcgg ctggcgtctt 60  
 accttggcgc tcttcttag agcttccact tacagttgtc aatttgaaga ggcagatgga 120  
 cagtgccctg catgatgcct cegtgttttag gcttattatg gacttgctca aagttgttct 180  
 gaataagtgc caggaaataa ggctctatc ttatttaggc ctctatcttc ggggtagctg 240  
 actaagtgtt gttatgtac ggaatccaga atctgtgaat tgaaatgaat gtaccgatg 299

<210> 897  
 <211> 454  
 <212> nucleic acid  
 <213> Zea mays

<400> 897

actttacttg gccngccagg tcaagatncc cgggtcgacc cagcntcca ccgcttcngc 60  
 cctcncgctt tcgttgtcat gaggagctc accgtggcaa gccgccaccc gggcgcgct 120  
 ttctctaccc gccgcgccc cttcttcac cccgccgag ccggaaggga ctgcacctc 180  
 cagcgttgct ggcgatggga aaaaactcag gacagctcct tcgggagctc actaaggacc 240  
 agccgccttc caagaactgt gcacggagat attttgaaga atttgctgc accaactgct 300  
 ggtgctgttt cagttgagca agctgaagct attgctgacc tccaaaagg tgaaatgtgg 360  
 tcggtgcaca agtttggtgg cacatgcatg gggacctctg agagaattca caatgttgct 420  
 gataaaancc ctninggattc cttcgaanaa ggaa 454

<210> 898

<211> 310  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 898  
  
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 cgctttctct acccgccgcc gcccccttct tcaccccgcc gcagccggaa aggactcgac 120  
 cttccagcgg tgctggcgat gggaaaaaac tcaggacagc tccttcggga antcactaag 180  
 gaccanccgc cttccaagaa ntgtgnacgg agatattttg aagnatttgc tcgcaccaac 240  
 tgctggtgct gtttcagttg agcaagctga agctattgct gacctccaa aaggtgacat 300  
 gtggtcggtg 310

<210> 899  
 <211> 276  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 899  
  
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 tctgctgcaa cgtagatcga tcattctcact gatagggat gtacagcgat cgtcattgat 120  
 tcttgaaaag gcattcaacg ttctgcgaag aaatgggtgtt aatgtccaga tgatctcaca 180  
 aggggcgtcc aaggtgaaca tctcttggt agtcaatgac agtgaggcaa aacagtgcgt 240  
 acaagccctc cattcagcgt tctttgagaa tggatc 276

<210> 900  
 <211> 240  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 900  
  
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 tcattctgctg caacgtagat cgatcatctc actgataggg aatgtacagc gatcgtcatt 120  
 gattcttgaa aaggcattca acgttctgag aagaaatggt gttaatgtcc agatgatctc 180  
 acaaggggag tccaaggtga acatattcctt ggtagtcaat gacagtgagg caaacagtg 240

<210> 901

<211> 279  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 901  
  
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 acagctccgc gttccggatg gngcccgang tgccgctcgt catccccgag gtcaacccccg 120  
 aggccatggc gaacgtccgc ctggggcagg gggcgattgt ggcaaaccg aattgctcga 180  
 ccatcatctg cntcatgggt gccacgccc tccatcgcca cgctaagggtg ttaaggatgg 240  
 ttgtcagcac ataccaagca gcaatgggtgc ggggtgctgc 279

<210> 902  
 <211> 254  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 902  
  
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 tgtgtggtga tcagatacgt aaagggtgccg cactcaatgc cgttcagatt gctgaaatgc 120  
 tgctgaagtg aatgcgactt aaccctcttg tccctccctc cctccctgtc cctaattgct 180  
 ctgatcaaatt gctggactgt actctgatta gtttgtcctc nattttgggtc gcctgttctg 240  
 tattctgcgg tgct 254

<210> 903  
 <211> 496  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 903  
  
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 caagatgttg tgtacacgga ttggatagat cgtccacaac ctcgattgga ccgtcataag 120  
 gaaatggggtt tacagtgacg ctgggaagag tgaggggaatg ttccgtgatg gacataagtt 180  
 ggtattgttt ggacataaca cagtggtagg agctgcgggt ggttcgatta ttantgaaag 240  
 ttgagtgtcg ccaaagggtt tattgcacaa aaaggaaaac aaagtgtata tgcggagtat 300  
 ctcgtcacta gtgtgtgtca ctagacagaa tagtgtgttt gtgttggttg tgtcgtgttt 360

tccaatgaan gacattgacc actagaacta tgtgaaaaaa aaaannagtc aacacaaaga 420  
 ntannaaaaa aggnocggccg ctctagagga tncaagctta cgtacccgtc atnacgtcan 480  
 aacccttcna aaatgt 496

<210> 904  
 <211> 435  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 904

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 cagtttgaaa agccacttga tgagcgcgta tgctcctcaa gtaggcctta atgagaaactc 120  
 aaagagagag ttctgggaag gcctggagga catgggttagt agtgtgccag ttggcgagaa 180  
 gctottcata ggaggagatc tcaatggcca tgatactgca agagaaattt tgagagcagc 240  
 tcctggtgtt accattattg atgaccgagc ttccaatcgc ttctctacac ctctggaggt 300  
 atcagacaaa gatgacgtan cantgggtag gattcgtcag gacttgtcct ggatggtaac 360  
 cganggtttg gacatatttg tgtgttggtg atcagatacg taaangcgcg ggactnaatg 420  
 ccgtcaaaat tggtt 435

<210> 905  
 <211> 287  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 905

atcaccagag tgaattagca gaagaattgg ctgaaaggaa gcatctcttc tgtgcacgcc 60  
 cacaaacatt gggagagacc atccgggcaa tggacctaga gacactgggtt cctatgagcc 120  
 aggggatgcc aaaccagtcg tcaccctgat caacagattt cttggcttcc agttgattga 180  
 cgaatctatt cggttgttct caatccatgt ggattagatg gtattgcgtc ggtttaattc 240  
 catagcaagt caaaatacat cctaaaccca ctcaatccac tccaatc 287

<210> 906  
 <211> 307  
 <212> nucleic acid  
 <213> Zea mays

<400> 906

ccgcgaccga tcgagcgaan ancccttccc gcgcccgcgc cgaaacccta gtcctcttta 60  
cgccatggcc accgtgtcgc tcaactccgc gccgggtcttc tccaccgagt ccggcggcgc 120  
cctgggcttc tgccaccatc ctccgcttcc cgccaaactt cgtacgccag ctacgacca 180  
aggcacgacg caactgcagc aacatcggcg tcgcgcagat cgtcgcgcgc gctggtccg 240  
actgcctcgc tgctcgcgc ctccgcgcg gccgatgtca gcgcaattcc taacgctaag 300  
gttgcg 307

<210> 907

<211> 265

<212> nucleic acid

<213> Zea mays

<400> 907

aaagtacatt gctggacaca acgatgttat tggaggatgc gtcagtggca gactgantgg 60  
tttccaaagt cntatattat caccatgttg ttggtggtgt ctaaccccga atgctncgta 120  
ctancctnnc ncnгааacct ncatctccgt gtgcgtgctg aataaacntg cttncggatg 180  
gccanttttt agaggaaacat ctaaagatgc cngctacta tcctggctgc tagccaaccc 240  
tgaacatcca ttgcccagag ncaat 265

<210> 908

<211> 267

<212> nucleic acid

<213> Zea mays

<400> 908

attttcagta aatttggtat cgaacacaca ctgaacatcc ctgcaacttt tgttggttaag 60  
angttgtttg tttaggacta taatctgtct agattatata atccaacaaa taatgtgtta 120  
tttgctggat tatataatct ggacagatta tattctcaaa caaacacccg ctaattcttc 180  
caggggacaa tcggagcgtg gaaacgatgc caaccaggac atactttaca accagaacgt 240  
ccaaagtgtg gtgccgtacc attcgag 267

<210> 909

<211> 96

<212> nucleic acid





<210> 912  
 <211> 497  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 912

ggtttttaan gggggggagg aagggaagc tttcccggtc cggattcccg ggtcgacca 60  
 cgcgtccoga gttaaaacat ctttttgng gtgggnaaat atggaaccta tttcangtta 120  
 nttggcctng nttggatttg ancccgtnng gtggagtaat ctaatactcc gcttgggatg 180  
 tcgatattgg agaggatgga attgaattgg ggttgaatac caaatcaggc atgatattga 240  
 aatgagatat aatttcaatt ctattgtttg gatgtcactg aattggagtt tgggaattatg 300  
 cgggtctaatt ctaggcaata ccgaggggggt gaggctttgt attggaagag ggggtttcta 360  
 gttatagtct aatttcaggg aattgggcct ctgattccaa atctcaattc cacatgcaac 420  
 gaatcaacat aatttangaa agctgattcc aattctcaat tctgtgattc aatctctaca 480  
 tccaaacggg gtataag 497

<210> 913  
 <211> 548  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 913

gnnnnngcga gagtttgttn gggngnnggg gggaaggang natntgctnn tccggtccgg 60  
 aattcccggg tcgaccacg cgtccgcgcg agcggcccc tttttttttt tttttttttt 120  
 ttttttttta ctaaaaccta tngnctgang ntcaggctaa tgacattttg tctttacaat 180  
 actgtgatgc aataaaactg ggcaaaccat tgataccacc gtatacagta actcaatacc 240  
 atccaattct atgaggatta aaaatgtatt ggaatggatt ggtatgtatt ttgacttgct 300  
 ataaatttaa accaactnaa taccatccaa ttcacttaga ttaacggtga aacgaacaag 360  
 gcctaagggg agtggttaaca gtctaacatc attgtacaac agcctttccg aagcccatag 420  
 acattgtgag acaagatccg taaaccatga atcgttcttc acttaggatg accatnttag 480  
 ctggtcacga aatggtaccg ggaatcacia atntttnaac gtcccaaang aacgatcctg 540  
 gangctgt 548

<210> 914  
 <211> 463  
 <212> nucleic acid  
 <213> Zea mays

<400> 914

cataaataag tcgtgttttg aagggttctn ntgaataaga taaaggctta gatacattna 60  
 gntgtatgaa cgggttnttt ttaggtgtat ctagtgtttg gagaaccgca acatgatgtc 120  
 tatgttgtac atgacctaag tgcatttctc taccaccagc actcttgccc aagcatggta 180  
 gaccatttgt gtttaatgtt cattacatgc taataaaaga tactggtggg aagggtaagg 240  
 ttatgtttta ggcacaccaa acaagaagga gtctaangaa attaagnnac caaggtctct 300  
 agtngagtaa gggggtagtc tttccattaa aattgactgg ctaaaattta agctcgggct 360  
 cttatgnng ttingngaac ctatgagaac cgacngnaag gnanttgggn aatcagtnag 420  
 tanttgaac naaacggtnt ggtaaagtga acccttttct cgt 463

<210> 915  
 <211> 267  
 <212> nucleic acid  
 <213> Zea mays

<400> 915

agctgaagga tttcttcaag cagtaggtgt gaagacctga agatcagctt tatctctctt 60  
 ttgttctctgc tagggccttc tcggttattc caattccatg tggattgaag tgtattaaaa 120  
 tggattttga cttgttatgg atttaaata acttaataacc actcaatcca catggattat 180  
 tggtgaaaca aacaagccct agtcgaaatt cttgttaagt tggtagtttg tcagaccoga 240  
 aatagagttt tttgggtttc gaatgga 267

<210> 916  
 <211> 284  
 <212> nucleic acid  
 <213> Zea mays

<400> 916

attttgaggc atggcagaca actgaggtcg aaatagcgtg gtagcttgga tctgcacatc 60  
 tatatgattg agggcctggg gtttaggatt ataatctgcc cagattatat aacctatctt 120

atTTTgaact aagtgttagt tcaaaataag ttagattata taatatggac aaattatatt 180  
 cccaatcaaa caggctatga ttggacttac tgatacccat tgagtagaac tggatcttgg 240  
 acagtntncg gggaaaacgg atggacctta attgaaaatt tgcg 284

<210> 917  
 <211> 247  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 917

ctgaggtcga aatagcgtgg tagcttggat ctgcacatct atatgattga gggcctggtg 60  
 tttaggatta taacctgcc agattatata acctatctta tttgaactaa gtgttagttc 120  
 aaaataagtt agattatata atctggacaa attatattcc caatcaaaca ggctatgatt 180  
 ggacttactg ataccattt gagtagaact gatcttgaca ggtatccgag aaaacgatga 240  
 ccttaat 247

<210> 918  
 <211> 254  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 918

caactggagg tcgaaatagc gtggttagctt ggatctgcac atctatatga ttganggcoct 60  
 ggtgttttagg attataatct gccagatta tataacctat cttattttga actaagtgtt 120  
 agttcanaat aagttagatt atataatatg gacaaattat attcccaatc aaacaggcta 180  
 tgattggact tactgatacc catttgagta gaactgatct tgacaggtat ccgagaaaac 240  
 gntgacctta attg 254

<210> 919  
 <211> 508  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 919

agangttttt nggntggggn ggnnganaag ttaccggtc cggattcccg ggtcgaccca 60  
 cgcgtccgct gggaagaagg atagccacgg atgctatcac cagcccggtg gtgaacacgt 120

cggcctactg gttcaacaac tcgcaagagc taatcgactt taaggagggg aggcattgcta 180  
 gcttcgagta tgggaggtat gggaacccga ccacggaggc attagagaag aagatgagcg 240  
 cactggagaa agcagagtcc accgtgtttg tggcgtcagg gatgtatgca gctgtggcta 300  
 tgctcagcgc acttgtccct gctggtgggc acattgtgac caccacggat tgctaccgca 360  
 agacaaggat ttacatggaa aatgagctcc ctaagagggg aatttcgatg actgtcatta 420  
 ggctgtctga catggatgct ctccaaaatg cgttggacaa caataatgta tctcttttct 480  
 tcacggagac tntacaaaat ccatttct 508

<210> 920  
 <211> 316  
 <212> nucleic acid  
 <213> Zea mays

<400> 920  
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 ccatagcaag ggagcgttgc tttgtatcga cagtactttt gcctccccta tcaatcagaa 120  
 ggcaactgact ttaggcgctg acctagtatt tcattctgca acaaagtaca ttgctggaca 180  
 caacgatgtt attggaggat gcgtcagtg cagagatgag ttggtttcca aagtccgtat 240  
 ttatcaccat gtggttggtg gtgttctaaa cccgaataac actgctcttc ggatggccca 300  
 gtttttagag gaacat 316

<210> 921  
 <211> 300  
 <212> nucleic acid  
 <213> Zea mays

<400> 921  
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 taatgtatct cttttcttca cggagactcc caaaaatcca tttctcagat gcattgatat 120  
 tgaacatgta tcaaatatgt gccatagcaa gggagcgttg ctttgtatcg acagtacttt 180  
 tgctcccct atcaatcaga aggcactgac ttaggcgct gacctagtta ttcattctgc 240  
 aacaaagtac attgctggac acaacgatgt tattggagga tgcgtcagtg gcagagatga 300

<210> 922

<211> 271  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 922  
  
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 tacatggaaa ctgagctccc caagagggga atttcgatga ctgtcattag gcctgctgac 120  
 atggatgctc tacaaaatgc gttggacaac aataatgtat ctcttttctt cacggagact 180  
 cccacaaatc catttctcag atgcattgat attgaacatg tatcaaatat gtgccatagc 240  
 aaggggagcgt tgctttgtat cgacagtact t 271

<210> 923  
 <211> 266  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 923  
  
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 gggagcgttg ctttgtatcg acagtacttt tgccctccct atcaatcaga aggcaactgac 120  
 tttaggcgtt gacctagtta ttcattctgc aacaaagtac attgctggac acaacgatgt 180  
 tattggagga tgcgtcagtg gcagagatga gttggtttcc aaagtccgta tttatcacca 240  
 tgtggttggt ggtgttctaa acccga 266

<210> 924  
 <211> 263  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 924  
  
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 aacaaggatt tacatggaaa ctgagctccc caagagggga atttcgatga ctgtcattag 120  
 gcctgctgac atggatgctc tacaaaatgc gttggacaac aataatgtat ctcttttctt 180  
 cacggagact cccacaaatc catttctcag atgcattgat attgaacatg tatcaaatat 240  
 gtgccatagc aaggggagcgt tgc 263

<210> 925

<211> 238  
 <212> nucleic acid  
 <213> Zea mays

<400> 925

gcgtaacctg ctcggtccg acgccagcct cgccgtccac gcgggggaga ggctgggaag 60  
 aaggatagcc acggatgcta tcaccacgcc ggtagtgaac acgtcggnct actggttcaa 120  
 caactcgcaa gagctaatacg actttaagga ggggaggcat gctagcttcg agtatgggag 180  
 gtatgggaac ccgaccacgg aggcattaga gaagaagatg agcgactgg agaaagca 238

<210> 926  
 <211> 332  
 <212> nucleic acid  
 <213> Zea mays

<400> 926

gccgcaactg cagcaacatc ggcgtcgctc agatcgctgc cgccgctgg tccgactgcc 60  
 ccgccgcctc cgcggcgcgg aggtcagcgc aattcccaac gctaagggtg cgcaaaccga 120  
 ccgcgctgt cttggccgag cgtaacctgc tcggctccga cgccagcctc gccgtccacg 180  
 cggggganag gctgggaaga aggatcgcca cggatgcgat caccacaccg gtagtgaaca 240  
 cgtcggccta ctggttcaac aactcgcaag agctaatcga ctttaaggag gggaggcatg 300  
 cttagcttca gtatgggagg tatgggaacc cg 332

<210> 927  
 <211> 226  
 <212> nucleic acid  
 <213> Zea mays

<400> 927

ctatgctcag tgcacttggt ccggctggtg ggcacattgt gaccaccacg gattgctacc 60  
 ggaaaacaag gatttacatg gaaactgagc tcccaagag gggaatttcg atgactgtca 120  
 ttaggcctgc tgacatggat gctctacaaa atgcgttga caacaataat gtancccttt 180  
 tcttcacgga gactcccaca aatccatttc tcagatgcat tgatat 226

<210> 928  
 <211> 228  
 <212> nucleic acid

<213> Zea mays

<400> 928

gatgactgtc attaggcctg ctgacatgga tgctctacaa aatgcgttgg acaacaataa 60  
tgtatctctt ttcttcaagg agactccac aaatccattt ctcagatgca ttgatattga 120  
acatgtatca aatatgtgcc atagcaaggg agcgttgctt tgtatcgaca gtacttttgc 180  
ctcccctatc aatcagangg catgacttta ggcgtgacc tagttatt 228

<210> 929

<211> 501

<212> nucleic acid

<213> Zea mays

<400> 929

gnnagnnaga ggttgatacg gaacgggggn ttaatggaat gctcgtaccg gtccggaatt 60  
cccgggtcga cccacgcgtc cgggttattg gaggatgcgt cagtggcaga gatgagttag 120  
tttccaaagt tcgnatttac caccatgtag ttggtggtgt tctaaacccg aatgctgcgt 180  
acottatcct tcgaggtatg aagacactgc atctccgtgt gcaatgtcag aacgacactg 240  
ctcttcggat ggcccagttt ttagaggagc atccaaagat tgctcgtgtc tactatcctg 300  
gcttgccaag tcaccctgaa catcacattg ccaagagtca aatgactggc tttggcggtg 360  
ttgttagttt tgaggttgct ggagactttg atgctacgag gaaattcatt gattctgtta 420  
aaatacccta tcatgcgcct tcttttgaag ctgtgagagc ataattgatc agcctgccat 480  
catgtcctac tgggattcaa a 501

<210> 930

<211> 489

<212> nucleic acid

<213> Zea mays

<400> 930

aacgttacgc cggncctag cctccaaaa ttcangggtc gacccacgcg tccggacaac 60  
ctgatcaggt tcagcattgg agtggaggat tttgaggacc ttaagaatga tcttgttcag 120  
gccctcgaga agatataagc tccaatccat ttgcattaat aaagttatga ggtgatgggt 180  
gtctggatct tgtagagatt tgtgatgata catgagctgt tgactgcgat taagttctct 240



tgtgcttatt ttattctttg aattcaataa gctgtggttg ctgcatcttg catgtagtgt 300  
 tggacttaat ttacttgctc aaattacaat agtttggtcc catgtacttg tgtccgcgcc 360  
 tccgcggttg tgccccaag tgtttttgtg tcatttgctg tgtaatgtga gattaagggc 420  
 acgatgtcct taatcaccat aataaatcca agccttaaat gtaaggattc tgggttgggg 480  
 atgaanaac 489

<210> 931  
 <211> 340  
 <212> nucleic acid  
 <213> Zea mays

<400> 931

tttaaaatac cctatcatgc gccttctttt ggaggctgtg agagcataat tgatcagcct 60  
 gccatcatgt cctactggga ttcaaaggag cagcgggaca tctacgggat caaggacaac 120  
 ctgatcaggt tcagcattgg tgtggaggat ttcgaggatc ttaagaacga tctcgtgcag 180  
 gccctogaga agatctaagc actctaata gtttgtattg acaaaatcat gaggtgatgg 240  
 ctgtcttggg tcttgtcaag atctgtgaca atgatatgag ctgatgactg cgaataagtt 300  
 ctcttttgcg tatttatccg tcaaatccat aagctgtgat 340

<210> 932  
 <211> 303  
 <212> nucleic acid  
 <213> Zea mays

<400> 932

ctgcatctcc gtgtgcaatg tcagaacgac actgctcttc ggatggccca gtttttagag 60  
 gagcatccaa agattgctcg tgtctactat cctggcttgc caagtcaccc tgaacatcac 120  
 attgccaaaga gtcaaagac tggctttggc ggtgttggtg gttttgaggt tgctggagac 180  
 tttgatgcta cgaggaaatt cattgattct gttaaaatac cctatcatgc gccttctttt 240  
 ggaggctgtg agagcataat tgatcagcct gccatcatgt cctactggga ttcaaaggag 300  
 cag 303

<210> 933  
 <211> 299  
 <212> nucleic acid

<213> Zea mays

<400> 933

tttacattta aggcattgat ttattatggt gattaaggac atcgtgcctt aatctcacac 60  
tacacagcaa atgacacaaa aacactaggg gcacaaccgc ggaggcgcg acacaagtac 120  
atgggaccaa actattgtaa ttgacaagt aaaattaagt ccaacactac atgcaagatg 180  
cagcaaccac agcttattga attcaaagaa taaaataagc acaagagaac ttaatcgag 240  
tcaacagctc atgtatcatc acaaattctc acaagatcca gacaaccatc acctcataa 299

<210> 934

<211> 345

<212> nucleic acid

<213> Zea mays

<400> 934

gggaacatct aaagattgcc cgtgtctact atcctggctt gcctagccac cctgaacatc 60  
acattgccaa gagtcaaagt actggctttg gtggtgtggt tagttttgag gttgctggag 120  
actttgatgc cacaaggaaa ttcattgatt ctgttaagat accctatcat gccccttctt 180  
tnggaggctg tgagagcata attgatcagc ctgctatcat gtcctactgg gattcaaagg 240  
agcagaggga catctatggg atcaaggaca acctgatcag gttcagcatt ggagtggagg 300  
ttttgaggac cttaagaatg atcttgttca ggccctcgag aagat 345

<210> 935

<211> 274

<212> nucleic acid

<213> Zea mays

<400> 935

agtcaaata ctggctttgg cgggtgtgtt agttttgagg ttgctggaga ctttgatgct 60  
acgaggaaat tcattgattc tgttaaaata ccctatcatg cgccttcttt tggaggctgt 120  
gagagcataa ttgatcagcc tgccatcatg tcctactggg attcaaagga gcagcgggac 180  
atctacggga tcaaggacaa cctgatcagg ttcagcattg gtgtggagga tttcgaggat 240  
cttaagaacg atctcgtgca ggccctcgag aaga 274

<210> 936

<211> 306  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 936  
  
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 tgcgtacctt atccttcgag gcatgaaaac acttcattctc cgtgtgcaat gtcagaataa 120  
 nactgctctt cggatggccc agttttttaga ggaacattcta aagattgccc gtgtctacta 180  
 tcctggcttg cctagccacc ctgaacatca cattgccaaag agtcaaata ga ctggctttgg 240  
 tgggtgtggtt agtttttnagg ttgctggaga ctttgatgcc acaaggaaat tcatcgattc 300  
 tgttaa 306

<210> 937  
 <211> 265  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 937  
  
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 gaagacactg catctccgtg tgcaatgtca gaacgacact gctcttcgga tggcccagtt 120  
 tttagaggag catccaaaga ttgctcgtgt ctactatcct ggcttgccaa gtcaccctga 180  
 acatcacatt gccaaagatc aaatgactgg ctttggcggg gttgttagtt ttgagggttc 240  
 tggagacttt gatgctacga ggaaa 265

<210> 938  
 <211> 322  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 938  
  
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 tcgattctgt taagatacnc tatcatgccc cttctttcgg aggctgtgag agcataattg 120  
 atcagcctgc tatcatgtcc tactgggatt caaaggagca gagggacatc tatgggatca 180  
 aggacaacct gatcaggttc agcattggag tggaggattt tgaggacctt aagaatgatc 240  
 ttgttcaggc cctcgagaag atataagctc caatccattt gcattaataa agttatgagg 300

tgatggttgt ctggatcttg ta

322

<210> 939  
<211> 283  
<212> nucleic acid  
<213> Zea mays

<400> 939

cattgccaaag agtcaaata ctggccttgg cgggtgttgtt agttttgagg ttgctggaga 60  
ctttgatgct tacgagggaa attcattgat tctgttaaaa taccctatca tgcgccttct 120  
tttgaggagct gtgagagcat aattgatcag cctgccatca tgtcctactg ggattcaaag 180  
gagcagcggg acatctacgg gatcaaggac aacctgatca gggtcagcat ggtgtggagg 240  
attcgaggat cttaagaacg atctcgtgca ggccctcgag aag 283

<210> 940  
<211> 282  
<212> nucleic acid  
<213> Zea mays

<400> 940

ctttagatgt tcctctaaag attgcccgtg tctactatcc tggcttgccct agccaccctg 60  
aacatcacat tgccaagagt caaatgactg gctttggtgg tgtgggttagt tttgagggtg 120  
ctggagactt tgatgccaca aggaaattca tcgattctgt taagataccc tatcatgccc 180  
cttctttcgg aggctgtgag agcataattg atcagcctgc tatcatgtcc tactgggatt 240  
caaaggagca gagggacatc tatgggatca aggacaacct ga 282

<210> 941  
<211> 315  
<212> nucleic acid  
<213> Zea mays

<400> 941

gggattcaaa ggagcagagg gacatctatg ggatcaagga caacctgatc aggttcagca 60  
ttggagtgga ggattttgag gaccttaaga atgatcttgt tcangccctc gagaagatat 120  
aagctccaat ccatttgcat taataaagtt atgangtgat gggtgtctgg atcttgtaga 180  
gatttgtgat gatacatnag cnttttactg ngaataagtt ctcttgngct taatttantic 240

tttgnattca ataantctgn ggttgcttgg aaaaaaaaaa aannntnnnn nnnnnntnnn 300  
 tttttnnnnt nnnnt 315

<210> 942  
 <211> 261  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 942

atttatcacc atgtggttgg tgggtgttcta aaccogaatg ctgcgtacct tatccttoga 60  
 ggcatgaaaa cacttcatct ccgtgtgcaa tgtcagaata aactgctct tcggatggcc 120  
 cagtttttag aggaacatct aaagattgcc cgtgtctact atcctggctt gcctagccac 180  
 cctgaacatc acattgcaa gagtcaaag actggctttg gtgggtgtgg tagttttgag 240  
 gttgctggag actttgatgc c 261

<210> 943  
 <211> 277  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 943

ttagggcgct gacctagtta ttcattctgc aacaaagtac attgctggnc acaacgatgt 60  
 tattggagga tgcgtcagt gcanagatga gttggnntcc naagncgtat ttatcaccat 120  
 gtggttgggt gtgtcctaaa cccgaatgct gcgtacctta tccttcgagg catgaaaaca 180  
 cttcatctcc gtgtgcaatg tnagaataac actgctcttc ggatggccca gtttttagag 240  
 gaacatctaa agattgccc tgtctactat cctggct 277

<210> 944  
 <211> 250  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 944

tgcgtcagt gacagatga gttagtttcc aaagttcgta tttaccacca ttagttgggt 60  
 ggtgttctaa acccgaatgc tgcgtacctt atccttcgag gtatgaagac actgcatctc 120  
 cgtgtgcaat gtcagaacga cactgctctt cggatggccc nngttttaga ggagnatcca 180

aagatttcgc gnggtcaagt atccgggctt gccaaagtcac cctgaacatc acattgccaa 240  
gagtcaaattg 250

<210> 945  
<211> 303  
<212> nucleic acid  
<213> Zea mays  
  
<400> 945

aggaaatcat cgattctgtt aagaaaccct atcatgcccc ttctttcgga ggctgtgaga 60  
gcataattga cagcctgcta tcatgtctac tgggattcaa aggagcagag gggcatctat 120  
gggatcaagg acaaccttga caggtcagca ttggagtgga ggattttgag gaccttaaga 180  
atgatcttgn tcaggccctc gagaagatat aagctccaat ccattttgat taataaaggt 240  
atgaggtgat ggggtggctgg atcttgagag attgngatga tncatgagct gttgctgcna 300  
tta 303

<210> 946  
<211> 239  
<212> nucleic acid  
<213> Zea mays  
  
<400> 946

ctatcatgcc ctttctttcg gaggtgtga gaggcataatt gatcagcctg ctatcatgtc 60  
ctactgggat tcaaaggagc agaggacat ctatgggatc aaggacaacc tgatcaggtt 120  
cagcattgga gtggaggatt ttgaggacct taagaatgat cttgttcagg ccctcgagaa 180  
gatataagct ccaatccatt tgcattaata aagttatgag gtgatggttg ctggatctt 239

<210> 947  
<211> 248  
<212> nucleic acid  
<213> Zea mays  
  
<400> 947

atttatcacc atgtggttgg tgggtgttcta aaccggaatg ctgcgtacct tatccttcga 60  
ggcatgaaaa cacttcatct ccgtgtgcaa tgcagaata aactgctct tcggatggcc 120  
cagtttttag aggaacatct aaagattgcc cgtgtctact atccnggctt gctagccac 180



<211> 178  
 <212> nucleic acid  
 <213> Zea mays

<400> 951

cacaacgatg ttattggagg atgcgtcagt ggcagagatg agttggtttc caaagtccgt 60  
 atttatcacc atgtgggttg tgggtgttcta aaccggaatg ctgcgtacct tatccttcga 120  
 ggcatgaaaa cacttcatct ccgtgtgcaa tgtcagaata aactgtctct tcggangg 178

<210> 952  
 <211> 308  
 <212> nucleic acid  
 <213> Zea mays

<400> 952

ancgaanacn gctccccgcg ccgcgcgcaa aaccctagct tctcctactc catggccact 60  
 gtctcgctca ccccgagggc tgtcttctcc acggagtcgg gtggcgccct ggctctgtct 120  
 accatcctcc gctttccgcc aaactttgtc cgccagctta gcaccaaggc acgccgcaac 180  
 tgcagcaaca tcggcgctgc gcagatcgtc gccgcgcggt ggtccgactg cncgcngcc 240  
 tccgcggcgc ggaggtcagc gcaattccca acgctaaggt tgcgcaaccg tccgcngtcg 300  
 tcttggcc 308

<210> 953  
 <211> 269  
 <212> nucleic acid  
 <213> Zea mays

<400> 953

gccacctccc cgaaagctgc cgcgaccgat cgagcgaagn ncggctcccc gcncccgcgc 60  
 ccaaaacctt agcttctctt actccatggc cactgtctcg ctacccccgc aggtgtctt 120  
 ctccacggan tccggtggcg cctggcctc tgctaccatc ctccgctttc cgccaaactt 180  
 tgtccgccag cttagcacca aggcacgccg caactgcagc aacatcggcg tcgcgcagat 240  
 cgtcgccgcc gctggtccg actgccccg 269

<210> 954  
 <211> 276  
 <212> nucleic acid



<213> Zea mays

<400> 954

ctgcaaaatg ctgaagggtc gggctctggca ccttttgact gctggctttg cttgagggga 60  
atcaaaacca tggctctgcg ggtggagaaa caacaggcta atgccagaa gattgctgaa 120  
ttcctggcgt ctaccccgag ggtcaagcaa gtaaaactacg ctgggcttcc tgaccatcct 180  
gggcgagctt tacactatct ccaggcnaag ggancgggct ctgttctcag ttttcccacc 240  
ggntaaatgg cccntcnaaa gnagggtggg ggaacc 276

<210> 955

<211> 440

<212> nucleic acid

<213> Zea mays

<400> 955

cgagagtaga cgatcagaac caccatata gtcctacca gcaagttgtg ccaagaaatg 60  
gaatagttgt aaaacgagta gatacaacga aaattagtaga tgtggtgtct gcaattggac 120  
cctccactag actgggtttgg ctcgaaagtc ccacgaaccc tcgtcagcaa attactgaca 180  
ttaagacaat ctgagagata gcgcattctc atgggtgctct tgttttggtt gacaacagca 240  
tcatgtctcc agtgcctctcc cgtcctatag aactgggagc tgatatcgtg atgcactcgg 300  
ctaccaaatt tataacggga catagtgate ttaagggctg ggaaatcttg cagtgaagg 360  
tganaatttg ggctaaaaga aggtaagggt ttctgcaaaa tgctgaaag ggncgggtcc 420  
gggnaccttt tgactggctg 440

<210> 956

<211> 283

<212> nucleic acid

<213> Zea mays

<400> 956

cgctgtggag accaccaagt acttcagcgt aacagtcagc ttggggagcg tgaagtcct 60  
catcagcctg ccgtgcttca tgtcccacgc atcaatccct gcctcgggtcc gcgaggagcg 120  
tggcctaacc gacgacctcg tccggatata ggtcggcatc gaggatgtcg aggacctcat 180  
cgccgatctg gaccgcgcgc tcagaactgg cccggtgtag acatcgccga tccttaggtc 240

atgtcaagct atcttttgat gattcattgg ttgactgctt gcg 283

<210> 957  
 <211> 340  
 <212> nucleic acid  
 <213> Zea mays

<400> 957

caagtacttc agcgtaacag tcagcttcgg gagcgtgaag tccctcatca gcntgccgtg 60  
 cttcatgten caagcatcaa tccctgcctc ggtcccgcan gagcgtggcc taaccgacga 120  
 cctcgtccgg atatcgggtcg gcatcgagga tgcgaggac ctcacgccc atctggaccg 180  
 cgcgctcaga actggcccgg tgtagacatc gccgatcctt aggtcatgtc aagctatctt 240  
 ttgatgattc attgggttgac tgcttgctg atgataataa tgggaatgtt gcttgatac 300  
 gccgctgcta naaaaaaact gngtcnagnn acngggnnnc 340

<210> 958  
 <211> 257  
 <212> nucleic acid  
 <213> Zea mays

<400> 958

gcattctcat ggtgctcttg ttttggttga caacagcatc atgtctccag tgctctccc 60  
 tcctatagaa ctgggagctg atactgtgat gcactcggct accaaattta tagcgggaca 120  
 tagtgatctt atggctggaa ttcttgcaat gaagggtgag agtttggtta aagaggtagg 180  
 gtttctgcaa aangctgaag ggtcgggtct ggcacctttt gactgctggc ttgcttgag 240  
 gggaatcaaa accatgg 257

<210> 959  
 <211> 521  
 <212> nucleic acid  
 <213> Zea mays

<400> 959

ggnnnagggg ctnngtntn tgacntntac ttnntggcat gccgtaccgg tccggaattc 60  
 ccgggtcgac ccacgcgtcc ggggtgtctgc aattggaccc tccactagac tggtttggct 120  
 cgaaagtccc acgaaccctc gtcagcaa at tactgacatt aagacaatct cagagatagc 180

acatttctcat ggtgctcttg ttttggttga caacagcatc atgtctccag tgctctcccg 240  
tcttatagaa ctgggagctg atatcgtgat gcactcggct accaaattta tagcgggaca 300  
tagtgatctt atggctggaa ttcttgcaagt gaaggggtgaa aagtttggtt aaagaggtan 360  
ggtttctgca aaatgctgaa ngggtcgggt ctggcacctt ttgactgntg gctttgcttg 420  
aagggaatca aaaccatggc tntgcnggtg gagaaacaac aggctaattgc ccagaagatt 480  
gctgaatttc tggggtnnta cccganggtc aaagcaagta a 521

<210> 960  
<211> 265  
<212> nucleic acid  
<213> Zea mays

<400> 960  
atcgtgatgc actcggctac caaatttata gcgggacata gtgatcttat ggctggaatt 60  
cttgcaagtga agggtagagag tttggctaaa gaggtagggt ttctgcaaaa tgctgaaggg 120  
tcgggtcttg caccttttga ctgctggctt tgcttgaggg gaatcaaaac catggctctg 180  
cggttgagga aacaacagggc taatgccagc aagattgctg aattcctggc gtctcaccog 240  
agggtaagc aagtaaacta cgctg 265

<210> 961  
<211> 148  
<212> nucleic acid  
<213> Zea mays

<400> 961  
atatcgtgat gcactcggct accaaattta tagcgggaca tagtgatctt atggctggaa 60  
ttcttgcaagt gaagggtagag agtttggtta aagaggtagg gtttctgcaa aatgctgaag 120  
ggtcgggtct ggcacctttt gactgctg 148

<210> 962  
<211> 453  
<212> nucleic acid  
<213> Zea mays

<400> 962  
ggctggggcg tcgctgacgg tgatcgagg cgctccggc ggctccgaac gagatctgag 60

cgctccgca gtctccgtgg aggccctgga ctccgtcgcc tccgattctg acttagagac 120  
gaaggagccc agtgtgtcga cgatgctgac gagcttcgag aactcgttcg acaagtatgg 180  
ggctctgagc acaccgctgt accagaccgc cacctttaag cagccttcag ctacagatta 240  
tggaacttat gattacacta gaagtggtaa cctactcgt gatgttctcc agagcctcat 300  
ggctaagctt gagaaagcag atcaagcatt ctgcttcacc agcgggatgg cggcggttagc 360  
tgcagtaaca caccttcttc angctggaca agaaatagtt gctggtgang acatatatgg 420  
tggttctgat cggctctactt ttgcaagttg tgc 453

<210> 963  
<211> 230  
<212> nucleic acid  
<213> Zea mays

<400> 963  
cagcgccctcc gcagtctccg tggaggccct ggactccatc gcctccnatt ctgacttaga 60  
gacgaaggag cccagtgtgt cgacgatgcn nacgagnttc gagaactcgt tcgacaagta 120  
tggggctctg agcacaccgg tgtaccagac cgccaccttt aagcagcctt cagntacaga 180  
ttatggaact tatgattaca ctagaagtgg taaccctact cgtgatgntc 230

<210> 964  
<211> 212  
<212> nucleic acid  
<213> Zea mays

<400> 964  
cogtgctgga caccaacatc ctctgggtga accccgactg cggantcaag acccgaaagt 60  
acacggaggt caagcccgcc ctgaccaaca tggctccgc tgctaactca ttgcaccca 120  
gctngccagc gccaatgagc ncttttttgc tttntcgttg ggaggnggcn tnttanttct 180  
nttttttnnaa nanagggccccc cccctttttt nt 212

<210> 965  
<211> 275  
<212> nucleic acid  
<213> Zea mays

<400> 965

ggnccaaaca ccaagttctc ctatgcttct cacaaggctg ttaacgaata taaggaggnt 60  
aaggngctcn gtgttganen cgtgcnagta nttgttngnc tagtnntgta gctgtngatc 120  
tcaaagcctg ccaagggatg ggagaacgga tncccnctgg cttgccctta ctaggacgat 180  
nctccnctgc tacaaggaag tgnntggenn aattnaaatg nagnnnnggg ttccaggatt 240  
canttttang agcccgattg tgnngcnca tgaat 275

<210> 966  
<211> 254  
<212> nucleic acid  
<213> Zea mays

<400> 966

caagacccta acatctctga gcagtgtgac tgcttatggt tttgatcttg tccgtgggtg 60  
ggcttgtcac gagtgtggt ttccctgctg gaaagtacct ctttgcctgn gttgtggatg 120  
gacgcaacat ctgggctgat gatcttgcta catcnctnag naccncaagt cnnntgantn 180  
tgnttttggg aagnaangct tgnngancaa ctgctgcgcg tnaggcacng cgcntgnnnc 240  
tngtaaanaa caat 254

<210> 967  
<211> 313  
<212> nucleic acid  
<213> Zea mays

<400> 967

cgatatggnt gagtacttcg gtgagcagct ctccggtttt gcgttcactg ccaattgctg 60  
ggtgcagtct atnggtcaag gtgcgtcaag cctccgatca tctacggtga cgtgagccgg 120  
cccaacccca tgaccgtttc tngtcgaaga cggcacagag catgacgtct cgcccaatga 180  
aggggatgct gactggncat cacatctcac tgtccttcgt cggattnaca gcgaggttgn 240  
gantgtacag angtcttgga tcaagaagga gngaggncct gagctgtgta tccgtaccaa 300  
cgcgagtnnt gaa 313

<210> 968  
<211> 313  
<212> nucleic acid  
<213> Zea mays

<400> 968

ggangtngag natnttgagg ctgntgcat tcaggcgatc cngattgatg aggcngntct 60  
aagggaagnn ttggcctgcg cangttatag catgcantct acctgcaactg cgctgtccnc 120  
tctatcngaa tcaccnctn cagantccan nacncaacc agatccacac acatatgtgc 180  
tagtnaaact ntaacnanat tatnttnga atcatcgaca tggacgcaga tgtgatcacc 240  
atcgagaact cccgtccgac gagaagctac tttccgtgtt ccgngagggn gtgaagtacg 300  
gagctgggat cng 313

<210> 969

<211> 222

<212> nucleic acid

<213> Zea mays

<400> 969

caaggccaaa aaagatctct gaggagggan tacgtcaactg ncatcaagga ggaaatcaag 60  
gnaggctcgtt aaggggnaag aagagcttga cattgatgtg cttgtgcatg gcnagcctga 120  
gagaaacgat atggttgagt acttcggtga gcaactctcg gtttttcntt nactgncaat 180  
ggctnnntnc aanntantng nnnaagggnnc ntnaagnctn na 222

<210> 970

<211> 534

<212> nucleic acid

<213> Zea mays

<400> 970

agntnntnnn tggnggggga aggtatgnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60  
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nncgngctcg acaccaacat nctctgggtg 120  
aaccctgact gtggtctcaa gacacgcacg tncacagagg tcaagcccg ctagaccaaca 180  
tggtctctgc accaagctca tncgcacna acttgnacg nngaaatgag gtctctngata 240  
gntccatggt ctgatagcnc ggaatgagcc agntgttttg aataatttgg gtgntacccc 300  
ctgnncnatg gnggaagtgn taggttaggc tctcattggn gggatacgcc gntacaagat 360  
gtgttttaag nnggagtggn gagnttttct ttgngntatg nntctgggng tatgtttnac 420  
gctttgntta taacagaagn tgaaatncgc anngttgcta tacttngcc aagtnaccaa 480

agctgaccng angaanaact gcagngcacc cttacacacc aangngggaa acgg 534

<210> 971  
 <211> 435  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 971

ccacgcgttc gcngacgcgt gcgggaacna ccngtctaag ttngagacat nctacnaant 60  
 nncnattgca atcaaaaagg aggttgagga tcttgaggct gatggtattc aggtgatcca 120  
 aatcnatngg gcanctgtaa taggagntc tggcactacn caagttacat natgcattct 180  
 acctgnaactg tncgtgccac tntttnaaga tcacnaactg nngatttnaa gacaccacnc 240  
 agatncacac cnacatgtgc tantcnaact tcaacnacat catatactcc attatcgaca 300  
 tggatgccga tntgatcacn atcgagaact cccggttntg actagaanct actgtccnga 360  
 ntttcgtgan ggtatgaagt acggagctgg cattgaacct ggtgtctact acattcatct 420  
 cctaggattc ccttc 435

<210> 972  
 <211> 430  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 972

cccactcttc cnnccactct tncgngnttt angacaaaact tgtcatatna ncttctctggg 60  
 ggctcattca caccgntgng gatcttctta acnanactaa tcttgacacc gaaattaant 120  
 cttgncttgc ttttgctgcc cagaaagctg tngaggttna tgctcttgct aatgcattgg 180  
 ctgntcaaaa agatnangct tacttcgcan caaatgcttc tgctcatcct cgannaaann 240  
 ctacccccgt gtgaccaatn aaaaantcca taangctnct gntgctntca aagctcttac 300  
 cacctccgtg ngaccaacat taactgctag attggatgcc ccatcaaaaa naatcttaac 360  
 cttcccatte cttccancac tacaatcggn tccatccaca aaccogttna acctcatnaa 420  
 gatccccngt 430

<210> 973  
 <211> 266  
 <212> nucleic acid

<213> Zea mays

<400> 973

ttcgacttca ggccccgggat gatcagcatc aacctcgacc tgnnnngaagn gcggcaacaa 60  
 nttcatcaag accgccgcct aacggccact tcggccgtga cgacgccgac nttcacctgg 120  
 gaggtggtga agccccctnaa gttcgacaag gcatcggntt aaggttggga gtgtcaactgt 180  
 ggncatgagg actancttcc tctggctctt ctgttacctg naagattgct gctgntggat 240  
 gagtgtgttt gatcatgact ggctgc 266

<210> 974

<211> 303

<212> nucleic acid

<213> Zea mays

<400> 974

gttcaactgcc aacggatggg tgcaatccta tggatcacgc tgtgtgaagc caccattat 60  
 ctacggtgat gtcagccggc cgaaccccat gactgttttc tggccaaga tggcacagag 120  
 catgaccct cgtcccatga agggaatgtt gactggtccg gtcacaatcc tcaactggtc 180  
 attcgtcagg aacgaccagc ctaggtttga gacatgctac caaatagctc ttgcaatcaa 240  
 aaaggagggtt gaggatcttg aggctgctgg tattcagggtg atccagatcg atgaggcagc 300  
 tct 303

<210> 975

<211> 296

<212> nucleic acid

<213> Zea mays

<400> 975

caacggatgg gtgcaatcct atggatcacg ctgtgtgaag ccaccatta tctatggtga 60  
 tgtcagccgg ccgaacccca tgactgtntt ctggccaag atggcacaga gcatgacccc 120  
 tcgtcccatg aagggaatgt tgactggtcc ggtcacaatc ctgaactggt cattcgtcag 180  
 gaacgaccag ctaggttttg agacatgcta ccaaatagct cttgcaatca aaaaggaggt 240  
 tgaggatctt gaggctgctg gtattcagggt gatccagatc gatgaggcag ctctaa 296

<210> 976



<211> 279  
<212> nucleic acid  
<213> Zea mays

<400> 976

cactgccaac ggatgggtgc aatcctatgg atcacgctgt gtgaagccac ccattatcta 60  
tggatgatgtc agccggccga accccatgac tgttttctgg tccaagatgg cacagagcat 120  
gacccctcgt cccatgaagg gaatgttgac tggtcgggtc acaatcctga actggtcatt 180  
cgtcaggaac gaccagccta ggtttgagac ntgctacaa ntagctcttg cantcaaaaa 240  
ggaggttgag gntcttgagg ctgctggtat tcaggtgat 279

<210> 977  
<211> 220  
<212> nucleic acid  
<213> Zea mays

<400> 977

ggaacggncc agnacgcgtg gngtncagc cggccgaacc ccatnactgt tttctggtgc 60  
aagacggcac agagcatgac cgctcgctcc atgaaggga tgttgactgg tccagtcaca 120  
atcctcaact ggtcatttgt caggaacgat cagcctaggt ttgagacatg ctaccaata 180  
gctcttgcaa taaaaaagga ggttgaggat cttgaggctg 220

<210> 978  
<211> 295  
<212> nucleic acid  
<213> Zea mays

<400> 978

cctgaacttg gcccaagcac caagttcaca tacgcttctc acaaggctgt ttctgagtac 60  
aaggaggcaa aggcgctcgg cattgataca gtcccagtcg ttgttgacc agtgatcac 120  
ttgctcctct ctaagcctgc caagggtgtg gagaaatctt tctctcttct ttnacttctt 180  
ggtagcattc ttcccatcta caaggagggt gttgctgagc tgaaggcagc tggtagcttca 240  
tggattcaac ttgatgagcc tacccttggt aaagaccttg atgctcncga attgg 295

<210> 979  
<211> 259  
<212> nucleic acid

<213> Zea mays

<400> 979

cttctcacaa ggctgtttct gagtacaagg aggcaaaggc gctcggcatt gatacagtcc 60  
cagtgtttgt tggaccagtg tcataacttgc tcctctctaa gcctgccaaag ggtgtggaga 120  
aatctttctc tcttctttca cttcttggta gcattcttcc catctacaag gaggttgttg 180  
ctgagctgaa ggcagctggg gcttcatgga ttcaacttga tgagcctacc cttgttaaag 240  
accttgatgc tcacgaatt 259

<210> 980

<211> 302

<212> nucleic acid

<213> Zea mays

<400> 980

gtttctgagt acaaggaggc aaaagcgctc ggtattgata ctgtcccagt gcttgttgga 60  
ccagtgtcat acttgctcct ctctaagcct gccaaagggtg tggagaaatc tttctctctt 120  
ctttcgcttc ttggcagcat tcttcccatc tacaaggagg ttgtttctga actgaaggca 180  
gctgggtgctt catggattca gtttgatgag cctacccttg ttaaagacct tgatgttcac 240  
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<210> 981

<211> 284

<212> nucleic acid

<213> Zea mays

<400> 981

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tcacgaattg gcagcattct cttcagcata tgctgaacta gagtcggcat tctctggatt 120  
gaatgtgctt atcgagacat actttgctga cattcctgct gagtcctaca agaccctcac 180  
gtcattgagt ggtgtgactg cttacgggtt tgatcttacc cgtggagcca agacccttga 240  
tcttatcagg agcagttccc ctctgggaag tacctcttcg ctgg 284

<210> 982

<211> 314  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 982  
  
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 agaccttgat gctcaogaat tggccgcatt ctcttcagca tatgctgaac tggagtcac 120  
 gttctctgga ttgaatgtgc ttatcgagac atacttcgct gacattcctg ctgagtccta 180  
 caagaccctc acatcaatga gtgggtgtga ctgcctacgg ntctgatctt aaccgtngag 240  
 ccaagaccct tgatottatc aggagcagct tcccctctgg gaagtacctc ttctgtgtgt 300  
 tgtagatgga cgca 314

64360  
 <210> 983  
 <211> 231  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 983  
  
 atacaaacta ccactttatt gtccctgaac ttggcccaag caccaagttc acatacgtt 60  
 ctcacaaggc tgtttctgag tacaaggagg caaaggcgct cggcattgat acagtcccag 120  
 tgcttggttg accagtgtca tacttgctcc tctctaagcc tgccaagggt gtggagaaat 180  
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<210> 984  
 <211> 229  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 984  
  
 gtttcngagt acaaggaggc aaaggcgctc ggcattgana cagtcccagt gncttgtnng 60  
 accagtgtca tacttgctcc tctctaagcc tgccaagggt gtggagaaat ctttcncnct 120  
 tctttcacnt cttggtagca tncntcccat ctacaaggag gttgtngcng agcngaaggc 180  
 agcnggtgct tcatggattc aacttgatga gcctaccctt gttaaagac 229

<210> 985  
 <211> 300  
 <212> nucleic acid

<213> Zea mays

<400> 985

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tgagctgaag gcagctggtg cttcatggat tcaacttgat gagcctaccc ttgttaaaga 120  
ccttgatgct cacggaattg gccgcattct cttcagcata tgctgaactg gagtcacgt 180  
tctctggatt gaatgtgctt atcgagacat acttcgctga cattcctgct gagtcacctac 240  
aagaccctca catcattgag tgggtgtgact gcttacgggt tcgatcttat ccgtggagcc 300

<210> 986

<211> 177

<212> nucleic acid

<213> Zea mays

<400> 986

gaaatctttc tctcttcttt cgcttcttgg cagcattctt cccatctaca aggagggttg 60  
ttctgaactg aaggcagctg gtgcttcatg gattcagttt gatgagccta cccttgtaa 120  
agaccttgat gttcacgaat tggcagcatt ctcttcagca tatgctgaac tagagtc 177

<210> 987

<211> 145

<212> nucleic acid

<213> Zea mays

<400> 987

ggaggntggt gctgagctga aggcagctgg tgcttcatgg attcagcttg atgagcctac 60  
ccttggttaa gaccttgatg ctacgaatt ggccgcattc tcttcagcat atgctgaact 120  
ggagtctngt tctctggatt gaatg 145

<210> 988

<211> 189

<212> nucleic acid

<213> Zea mays

<400> 988

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tcttcagcat atnctgaact ggagtcacg ttctcnggat tgaatgtgct atcgagacan 120

acttcgctga catnctgctg agtcctacaa gaccctcaca tcatgagtgg tgtgatnctt 180  
acggtttctg 189

<210> 989  
<211> 532  
<212> nucleic acid  
<213> Zea mays  
  
<400> 989

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gattccccggg tcgacccaag cgtccgcgga cgcgtgggta caggcttact tcgcagcaaa 120  
tgctgctgct caggcctcga ggaaatcctc accccgtgtg accaatgaag aagtccagaa 180  
ggctgctgct gctctcaagg gctctgacca ccgcgtgcg accaacgtta gtgctagatt 240  
ggatgcccag cagaagaage ttaacctccc catccttccc accactacaa tcggttcggt 300  
cccacaaacc gttgagctca ggagggtccg ccgtgagtac aaggcaaaaa agatctctga 360  
ggaggaatac gtcactgcc acaaggagga aatcaacaag gtcgttaagc tccaagaaga 420  
gcttgacatt gatgtgcttg tgcatggcga gcttgagaga aacgatattg ttgagtactt 480  
cggtgagcag ctctccgggtt ttgcgttcac tgccaatggc tgggtgcagt ct 532

<210> 990  
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aacgttagtg ctagattgga tgcccagcag aagaagctta acctccccat ccttcccacc 180  
actacaatcg gttcgttccc acaaaccgtt gagctcagga gggtcgcgcg tgagtacaag 240  
gcaaaaaaga tctctgagga ggaatacgtc actgccatca aggaggaaat caacaaggtc 300  
gttaagcttc aagaagagct tgacattgat gtgcttgtgc atggcgagcc tgagagaaac 360  
gatatgggtg agtacttcgg tgagcagctc ttcggttttg cgttcactgg caatggctgg 420  
gtncagtctt atgggtcaag g 441

<210> 991  
 <211> 524  
 <212> nucleic acid  
 <213> Zea mays

<400> 991

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 tactccaact tcaacgacat catncactng atcatcgaca tggacgcgga cgtgatcacc 180  
 atcgagaact cgcggtccga cgagaagctg ctctccgtgt tccgtgaggg tgtcaagtac 240  
 ggcgcgggca tcggccccgg tgtctacgac atccactccc ccaggatccc gtgcggcggag 300  
 gagatcgccg accgcatcga caagatgctg gccgtgctgg acaccaacat cctctgggtg 360  
 aaccccgact gcggcctcaa gaccgcgaag tacacggagg tcaagccgc cctgaccaac 420  
 atggtctccg ctgctaagct cattcgcacc cagctcgcca agcgccaaag tgagcacctt 480  
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<210> 992  
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 <212> nucleic acid  
 <213> Zea mays

<400> 992

gaattnccaa ctgcgatttc caggacacca cccagatcca caccacntg tgctactcca 60  
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 actcgcggtc cgacgagaag ctgctctccg tgttccgcga ggcgtcaag tacggcgcgg 180  
 gcatcgcccc cgggtgtctac gacatccact ccccaggat cccgtcggct gaggagatcg 240  
 ccgaccgcat cgacaagatg ctggcgggtgc tggacaccaa catcctctgg gtgaaccccg 300  
 actgcggcct taagaccgc aagtacacgg aggtcaaagc ccgcctgacc aacatggtct 360  
 ncgncgctaa gctcatncgc acccagntcg ccancgcaa gtgancacct ttttttttgg 420  
 cctttttngn ttccga 436

<210> 993  
 <211> 330  
 <212> nucleic acid

<213> Zea mays

<400> 993

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catcagtgcc atcaaggaag aaatcagcaa ggttgtcaag atccaagagg agcttgacat 180  
tgatgtgctt gtgcatggag agccagaaaag aaatgacatg gttgagtact tcggtgagca 240  
attatctggt tttgcattca ctgccaacgg atgggtgcaa tcttatggat cacgttgtgt 300  
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<210> 994

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<212> nucleic acid

<213> Zea mays

<400> 994

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aaggaagtgc aggatcttga ggctgggtgt attcagggtta tccaaatcga cgaggctgca 180  
ctgagagaag ggctgccgct ccgcaaggct gaacatgcat tctacttga ctgggctgcc 240  
actccttcag aattaccaac tgcgagatcc angacaccac ccagatccac acccacatgt 300  
gtacttcaa cttcaacgac atcatncact cgatcatcga catggacgcc gacgtgaaca 360  
ccattgagaa cttgcgggcc gacnagaaac ttntnttccg ngttncnca aggccgtnaa 420  
gtccgggncc gggcatnt 438

<210> 995

<211> 469

<212> nucleic acid

<213> Zea mays

<400> 995

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caatcaaaaa ggaggttgag gatcttgagg ctgctggtat tcaggatgat cagatcgatg 180

aggcagctct aagggagggg ctgccactac gcaagtcaga gcatgcattc tacctggact 240  
 gggctgtcca ctctttcagg atcaccaact gcggagtcca ggacaccacc cagatccaca 300  
 cccacatgtg ctactccaac ttcaacgaca tcatccactc catcatcgac atggatgccc 360  
 gatgtgatca cgatcgagaa ctcccgggtc gaccagaagc tactgtccgt cttnccgtgag 420  
 ggtgtgaaat accggaactg gcattggccc tgggtgtctac gacattcat 469

<210> 996  
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 cgatgtgatc acgatcgaga actcccggtc tgacgagaag ctactgtccg tcttccgtga 180  
 ggggtgtgaag tacggagctg gcattggccc tgggtgtctac gacatccact ctcttaggat 240  
 tccctccacc gaggagatcg cagaccgctg cgagaagatg ctgcgccgtg tcgacaccaa 300  
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<210> 997  
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 <212> nucleic acid  
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<400> 997

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 gagcctgaga gaaacgatat ggttgagtac ttcggtgagc agctctccgg ttttgcggtc 120  
 actgccaatg gctgggtgca gtcttatggg tcaagggtcg tcaagcctcc gatcatctac 180  
 ggtgaccgtg agccgcccc accccatgac cgttttctgg tcgaagacgg cacagagcat 240  
 gacgtctcgc ccaatgaagg ggatgctgac tggcccagtc acaatcctca actggtcctt 300  
 cgtccggaat gaccagccga ggtttgagac ttgctaccag atccgctctt gcgatcaaga 360  
 aggaaagtcg aggatcttga agctggtggg tattcaaggg tatccaaatc gaccagggct 420  
 tgcactnaaa gaaanggctt nccgnttcgn aa 452



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<400> 998

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 aagctgctct ccgtgttccg tgaggggtgc aagtacggcg cgggcatcgg ccccggtgtc 180  
 tacgacatcc actccccag gatcccgctg gccgaggaga tcgccgaccg catcgacaag 240  
 atgctggccg tgctggacac caacatcctc tgggtgaacc ccgactgcgg cctcaagacc 300  
 cgcaagtaca cggaggtcaa gcccgccctg accaacaatg tctccgctgc taagtcatt 360  
 cgcaccagc tcgccagcgc caagtgagca ctttttttg cttttttcgt ttccgaagaa 420  
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<400> 999

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 catccactcg atcatcgaca tggacgcgga cgtgatcacc attgagaact cgcggtccga 180  
 cgagaagctg ctctccgtgt tccgcgaggg cgtcaagtac ggcgcgggca tcggccccgg 240  
 tgtctacgac atccactccc ccaggatccc gtcgggtgag gagatcgccg accgcatcga 300  
 caagatgtgg cgggtgtggac accaacaatc tctgggtgaa cccgattgcg gct 353

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 <212> nucleic acid  
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<400> 1000

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 tacctggact gggctgtcca ctctttcagg atcaccaact gcgaggtcca ggacaccacc 120  
 cagatccaca cccacatgtg ctactccaac ttcaacgaca tcatccactc catcatcgac 180  
 atggatgccg atgtgatcac gatcgagaac tcccggctcg acgagaagct actgtccgtc 240  
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<210> 1001  
 <211> 347  
 <212> nucleic acid  
 <213> Zea mays

<400> 1001

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 ccacaaccac aattgnntca ttccctcaga ctgtggaact caggagggtc cgctcgtgagt 240  
 acaaggcaaa gaagatctcc gaggaggaat acatcagtgc catcaaggaa gaaatcagca 300  
 aggttgtcaa gatccaagag gagcttgaca ttgatgtgct tgtgcat 347

<210> 1002  
 <211> 319  
 <212> nucleic acid  
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<400> 1002

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 ggttgtcaag atccaagagg agcttgacat tgatgtgctt gtgcatggag agccagaaag 120  
 aaatgacatg gttgagtact tcggtgagca attatctggt ttgcatcca ctgccaacgg 180  
 atgggtgcaa tottatggat cacgttgtgt gaagccaccc attatctatg gcgatgtcag 240  
 cggccgaac cccatgactg ttttctggtc caagacggca cagagcatga ccgctcgtcc 300  
 catgaaggga atgttgact 319

<210> 1003

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 gactaagctt gacagcgaga ttaagtcttg gcttgctttt gctgcccaga aggttggtga 120  
 gggttgatgct cttgctaagg cattggctgg tcaaaaggat gaggttact tcgcagcaaa 180  
 tgctgctgct caggcctcga ggaaatctc accccgtgtg accaatgaag aagtccagaa 240  
 ggctgctgct gctctcaagg gctctgacca ccgccgtggg accaacgtta gtgctagatt 300  
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gaagcgggacg tgatcaccat cgagaactcg cgggccgacg agaagctgct ctccgtgttc 240  
 cgtgaggggtg tcaagtacgg cgcgggcatc ggccccgggtg tctacgacat ccactccccc 300  
 ag 302

<210> 1006  
 <211> 296  
 <212> nucleic acid  
 <213> Zea mays

<400> 1006

gcagaaaaag ctcaaccttc ctgtccttcc cacaaccaca attggttcat tccctcagac 60  
 tgtggaactc aggaggggtcc gtcgtgagta caaggcaaag aagatctccg aggaggaata 120  
 catcagtgcc atcaaggaag aaatcagcaa ggttgtcaag atccaagagg agcttgacat 180  
 tgatgtgctt gtgcatggag agccagaaaag aaatgacatg gttgagtact tcggtgagca 240  
 attatctggt tttgcattca ctgccaacgg atgggtgcaa tcttatggat cacgtt 296

<210> 1007  
 <211> 537  
 <212> nucleic acid  
 <213> Zea mays

<400> 1007

gntngagtcn ggcttcggcn ttgatgatng cgtantcntt cngnangccg ntngaattc 60  
 ccgggtcgac ccacgcgtcc ggnaggaata cgtcactgcc atcaaggagg aaattaacaa 120  
 ggtcgttaag cttcaagaag agcttgacat tgatgtgctt gtgcatggng agcctgagag 180  
 aaacgatatg gttgagtact tcggtgagca gctctncggt tttgcgttca ctgccaatgg 240  
 ctgggtgcag tcttatgggt caaggtgcgt caagcctccg atcatctacg gtgacgtgag 300  
 ccgtcccaac cccatgaccg ttttctggtc gaagacggcn canagcatga cgtctcgccc 360  
 aatgaaaggg gatgctgact ngcccagtca caatcctcaa ctggtncctt gtcccggaat 420  
 naccagccca gggtttgaga nttgctacca gaatcggctt cttgcgatca agaaaggga 480  
 gtncaaggga tctttgangn tgggtggtn ttaangttta ttncaaaatc gacnaag 537

<210> 1008  
 <211> 347  
 <212> nucleic acid

<213> Zea mays

<400> 1008

cgttttcttg tcgaagacgg ggcagagca tgacgtctcg cccaatgaag gggatgctga 60  
ctggcccagt cacaatcctc aactggtcct tcgtccggaa cgaccagccg aggtttgaga 120  
cttgctacca gatcgctctt gcgatcaaga aggaagtcga ggatcttgag gctgggtgga 180  
ttcaggttat ccaaatcgac gaggtgcac tgagagaagg gctgccgctc cgcaaggctg 240  
agcatgcatt ctacttgga c tgggctgtcc actccttcag aattaccaac tgcgagatnc 300  
aggacaccac cagatccaca cccacatgtg ctactccaac ttcaacg 347

<210> 1009

<211> 339

<212> nucleic acid

<213> Zea mays

<400> 1009

gaggatcttg aggctgctgg tattcagggtg atccagattg atgaggcagc tctaagggag 60  
ggtctgccac tgcgcaagtc agagcatgca ttctacctgg actgggctgt ccaactcttc 120  
agaatcacca actgcgaggt ccaggacacc acccagatcc acacccatat gtgctactcc 180  
aacttcaacg acatcatcca ctccatcatc gacatggacg cagatgtgat caccatcgag 240  
aactcccggg cgcagagaaa gctactgtcc gtgttccgag aggtgtgaag tacggagctg 300  
gaatcgggtc agtgtctagg aatccactcg ccaagtc 339

<210> 1010

<211> 313

<212> nucleic acid

<213> Zea mays

<400> 1010

ccgtccgca aggtgagca tgcattctac ttggactggg ctgtccactc cttcagaaac 60  
accaactgag agatccagga caccaccag atccacaccc acatgtgcta ctccaacttc 120  
aacgacatca tccaatcgat catcgacatg gacgaggacg tgatcaccat cgagaactcg 180  
cgggccgacg agaagctgct ctccgtgttc cgtgagggtg tcaagtacgg cgcgggcatc 240  
ggccccgggt tctacgacat ccaactcccc aggatcccggt cggccganga gatcgccgac 300

cgcatcgaca aga

313

<210> 1011  
<211> 310  
<212> nucleic acid  
<213> Zea mays

<400> 1011

caacaaggtc gttaagcttc aagaagagct tgacattgat gtgcttgtgc atggcgagcc 60  
tgagagaaaac gatatggttg agtacttcgg tgagcagctc tccggttttg cgttcactgc 120  
caatggctgg gtgcagtctt atgggtcaag gtgcgtcaag cctccgatca tctacggtga 180  
cgtgagccgc cccaacccca tgaccgtttt ctggtcgaag acggcacaga gcatgacgtc 240  
tcgccccaatg aaggggatgc tgactggccc agtcacaatc ctcaactggt ccttcgtccg 300  
gaatgaccag 310

<210> 1012  
<211> 319  
<212> nucleic acid  
<213> Zea mays

<400> 1012

acgatatggt tgagtacttc ggtgagcagc tctccggttt tgcgttcact gccaatggct 60  
gggtgcagtc ttatgggtca aggtgcgtca agcctccgat catctacggt gacgtgagcc 120  
gccccaaacc catgaccgtt ttctgggtcga agacggcgca gagcatgacg tctcgcccaa 180  
tgaaggggat gctgactggc ccagtcacaa tctcaactg gtccttcgtc cggaatgacc 240  
agccgaggtt tgagaattgc taccagatcg ctcttgcat caagaaggaa gtcgaggatc 300  
tcgaggctgg tggatttca 319

<210> 1013  
<211> 315  
<212> nucleic acid  
<213> Zea mays

<400> 1013

ctagattgga tgcccagcag aagaagctta acctcccat ccttcccacc actacaatcg 60  
gttcgttccc acaaaccgtt gagctcagga gggtcgccg tgagtacaag gcaaaaaaga 120

tctctgagga ggaatacgtc actgccatca aggaggaaat caacaaggtc gttaagcttc 180  
aagaagagct tgacattgat gtgcttgtgc atggcgagcc tgagagaaac gatatggttg 240  
agtacttcgg tgagcagctc tccggttttg cgttcaactgc caatggctgg gtgcagtctt 300  
atgggtcaag gtgcg 315

<210> 1014  
<211> 505  
<212> nucleic acid  
<213> Zea mays  
<400> 1014

aggggtgtnag gggggggaag aaaggaactg tcgtaccggt ccggaattcc cgggtcgacc 60  
cacgcgtccg cttgtccgtg gaacccaaac tcttgggctt gtcacgagtg ctggtttccc 120  
tgctggaaag tacctctttg ctgggtgttg ggatggacgc aacatctggg ctgatgatct 180  
tgctacatct ctcagcactc tccagtctct tgaggctgtt gttgggaagg acaagcttgt 240  
cgtatcaact tctgtctgc tcatgcacac cgctgtggat cttgtgaacg agactaagct 300  
tgacagcgag attaagtctt ggcttgcttt tgctgccag aaggttgttg aggttgatgc 360  
tcttgctaag gcattggctg gtcaaaagga tgaggcttac ttcgcagcaa atgctgctgc 420  
tcaagcctcg aggaaatcct naccctgtgt gaccaatgaa aaatccagaa ngctgctgct 480  
gntctttnaag gcttttgacc accgc 505

<210> 1015  
<211> 298  
<212> nucleic acid  
<213> Zea mays  
<400> 1015

gccgtgaata caaggcaaag aagatcaccg aggacgaata catcagtgcc atcaaggaag 60  
aaatcagcaa ggttgtcaag atccaagagg agcttgacat tgatgtgctt gtgcatggag 120  
agccagagag aaatgacatg gttgagtact tcggtgagca attatctggt ttgctgttca 180  
ctgccaacgg atgggtgcaa tcctatggat cacgctgtgt gaagccaccc attatctatg 240  
gtgatgtcag ccggccgaac cccatgactg ttttctggtc caagatggca cagagcat 298

<210> 1016





tgtggatctt gtgaacgaga ctaagcttga cagcgagatt aagtcttggc ttgcttttgc 120  
 tgcccagaag gttgttgagg ttgatgctct tgctaaggca ttggctgggc aaaaggatga 180  
 ggcttacttc gcagcaaag ctgcagctca ggctcgagg aaatcctcac cccgtgtgac 240  
 caatgaagaa gtccagaagg ctgctgctgc tctcaagggc tctgaccacc gccgtgagac 300  
 caacgttagt gctagattng atgccagca gaagaagcct aacc 344

<210> 1019  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays

<400> 1019

ccgccgtgct accactgttg ctgctagatt ggatgctcag cagaaaaagc tcaaccttcc 60  
 tgtccttccc acaaccacaa ttggttcatt cctcagact gtggaactca ggaggggtccg 120  
 tcgtgagtac aaggcaaaga agatctccga ggaggaatac atcagtgcc tcaaggaaga 180  
 aatcagcaag gttgtcaaga tccaagagga gcttgacatt gatgtgcttg tgcattggaga 240  
 gccagaaaga aatgacatgg ttgagtactt cggtagagca ttatctggtt tt 292

<210> 1020  
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 <212> nucleic acid  
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<400> 1020

accagccgag gtttgagact tgctaccaga tcgctcttgc gatcaagaag gaagtcgagg 60  
 atctcgaggc tgggtgtatt caggttatcc aaatcgacga ggctgcactg agagaagggc 120  
 tgccgctccg caaggctgag catgcattct acttgactg ggctgtccac tccttcagaa 180  
 tcaccaactg cgagatccag gacaccaccc agatccacac ccacatgtgc tactccaact 240  
 tcaacgacat catccactcg atcatcgaca tggacgcgga cgtgatcacc atcgagaact 300  
 cgcggtccga cg 312

<210> 1021  
 <211> 286  
 <212> nucleic acid  
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<400> 1021

caagatccaa gaggagcttg acattgatgt gcttgtgcat ggagagccag agagaaatga 60  
catggttgag tacttcggtg agcaattatc tggttttgcg ttcaactgcc aaggatgggt 120  
gcaatcctat ggatcacgct gtgtgaagcc acccattatc tacggtgatg tcagccggcc 180  
gaaccccatg actgttttct ggtccaagat ggcacagagc atgacccctc gtcccatgaa 240  
gggaatggtg actgggtccg tcacaatcct caactgggtca ttcgtc 286

<210> 1022

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 1022

atgtgctact ccaacttcaa cgacatcatc cactccatca tcgacatgga tgccgatgtg 60  
atcacgatcg agaactcccg atctgacgag aagctactgt ccgtcttccg tgaggggtgtg 120  
aagtacggag ctggcattgg ccttggtgtc tacgacatcc actctcctag gattccctcc 180  
accgaggaga tcgcagaccg cgtcgagaag atgctcgccg tgctggacac caacatcctc 240  
tgggtgaacc ctgactgtgg tctcaagaca cgcaagtaca cggaggt 287

<210> 1023

<211> 295

<212> nucleic acid

<213> Zea mays

<400> 1023

caaggntgtc aagatccaag aggagcttga cattgatgtg cttgtgcatg gagagccaga 60  
gagaaatgac atggttgagt acttcggtga gnaattatct ggttttgagt tcaactgcca 120  
cggatgggtg caatcctatg gatcacgctg tgtgaagcca cccattatct atggtgatgt 180  
cagccggccg aaccccatga ctgttttctg gtccaagatg gcacagagca tgacccctcg 240  
tcccatgaag ggaatgttga ctgggtccggt cacaatcctg aactgggtcat tcgtc 295

<210> 1024

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1024

agaagatcac cgatgacgaa tacatcagtg ccatcaagga agaaatcagc aaggttggtca 60  
agatccaaga ggagcttgac attgatgtgc ttgtgcatgg agagccagag agaaatgaca 120  
tggttgagta cttcgggtgag caattatctg gttttgcgtt cactgccaac ggatgggtgc 180  
aatcctatgg atcacgtgt gtgaagccac ccattatcta tggatgatgtc agccggccga 240  
accccatgac tgttttctgg tccaagatgg cacagagcat gacccctcgt cccatgaag 299

<210> 1025

<211> 312

<212> nucleic acid

<213> Zea mays

<400> 1025

tgagtacaag gcaaaaaaga tctctgagga ggaatacgtc actgccatca aggaggaaat 60  
caacaaggtc gttaagcttc aagaagagct tgacattgat gtgcttggtc atggcgagcc 120  
tgagagaaac gatatggttg agtacttcgg tgagcagctc tccggttttg cgttcactgc 180  
caatggctgg gtgcagtctt atgggtcaag gtgcgtcaag cctccgatca tctacggtga 240  
cgtgagccgc cccaacccca tgaccgtttt ctgggtcgaag aggcacagag catgacgtct 300  
cgcccaatga ag 312

<210> 1026

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1026

aaagatctct gaggaggaat acgtcactgc catcaaggag gaaatcaaca aggtcgttaa 60  
gcttcaagaa gagcttgaca ttgatgtgct tgtgcatggc gagcctgaga gaaacgatat 120  
ggttgagtac ttcggtgagc agctctccgg ttttgcgttc actgccaatg gctgggtgca 180  
gtcttatggg tcaaggtgcg tcaagcctcc gatcatctac ggtgacgtga gccgccccaa 240  
cccatgacc gttttctggt cgaagacggc acagagcatg acgtctcgcc caatgaagg 299

<210> 1027

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 1027

gtgtgaccaa tgaagaagtc cagaaggctg ctgctgctct caagggtctt gaccaccgcc 60  
gtgggaccaa cgttagtgt agattggatg cccagcagaa gaagcttaac ctccccatcc 120  
ttcccaccac tacaatcggg tegtccccc aaaccgttga gtcaggagg gtccgccgtg 180  
agtacaaggc aaaaaagatc tctgaggagg aatacgtcac tgccatcaag gaggaaatca 240  
acaaggctgt taagcttcaa gaagagcttg acattgatgt gcttgtgcat ggcgagcctg 300  
agagaaacgg 310

<210> 1028

<211> 305

<212> nucleic acid

<213> Zea mays

<400> 1028

gtctcgccca atgaagggga tgctgactgg cccagtcaca atcctcaact ggtccttcgt 60  
ccggaatgac cagccgaggt ttgagacttg ctaccagatc gctcttgcca tcaagaagga 120  
agtccaggat cttgaggctg gtggtattca ggttatccaa atcgacgagg ctgcactgag 180  
agaagggctg ccgctccgca aggctgagca tgcattctac ttggactggg ctgtccactc 240  
cttcagaatt accaactgcg agatccagga caccaccag atccacacc acatgtgcta 300  
ctcca 305

<210> 1029

<211> 330

<212> nucleic acid

<213> Zea mays

<400> 1029

cgctgaggat ctggagaagg ttgctaccga cctcagggcc agtatctgga agcagatggc 60  
tgatgctggg atcaagtaca tccccagcaa caccttctcg tactatgacc agntccttga 120  
caccactgcc atgctcggcg ctgtccccga acgctactca tggactggag gagagattgg 180  
gtttgacacc tactttctcca tggccagggg caacgccact gtccctgcta tggagannac 240  
caagtggttt gacaccaact accactttat tgtccctgaa ttgggcccac acaccaagtt 300

ctcctatgct tctcacaagg ctgttaacga

330

<210> 1030

<211> 269

<212> nucleic acid

<213> Zea mays

<400> 1030

gcattctacc tggactgggc tgtccactct ttcagaatca ccaactgagg agtccaggac 60

accacccaga tccacaccca tatgtgctac tccaacttca acgacatcat ccactccatc 120

atcgacatgg acgcagatgt gatcaccatc gagaactccc ggtccgacga gaagctactg 180

tccgtgttcc gcgaggggtgt gaagtacgga gctggaatcg gtccaggtgt ctacgacatc 240

cactcgccca ggatcccctc caccgagga 269

<210> 1031

<211> 311

<212> nucleic acid

<213> Zea mays

<400> 1031

ctcgcccaat gaaggggatg ctgactggcc cagtcacaat cctcaactgg tccttcgtcc 60

ggaatgacca gccgaggttt gagacttgct accagatcgt cttgcgatca agaaggaagt 120

cgaggatctc gaggctggtg gtattcaggt tatccaaatc gacgaggctg cactgagaga 180

agggctgccg ctccgcaagg ctgagcatgc attctacttg gactgggctg tccactcctt 240

cagaatcacc aactgcgaga tccaggacac caccagatc cacaccaca tgtgctactc 300

caacttcaac g 311

<210> 1032

<211> 382

<212> nucleic acid

<213> Zea mays

<400> 1032

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tcgatcatcg acatggacgc ggacgtgatc accattgaga actcgcggtc cgacgagaag 120

ctgctctccg tgntccgcga gggcgtcaag tacggcgcgg gcatcggcnc cggtgtctac 180

gacatccact cccccaggat cncgtcggct gangagatcg ccgaccgcat cgacaagatg 240  
 ctggcgggtgc tggacancaa catcctctgg gtgaaccccg actgcngcct caagacccgc 300  
 aagtacacgg agtcaagccc gcntgancaa catggtctcc gccgtaagct catccgnacc 360  
 agctcgccag cgcaatgana ac 382

<210> 1033  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1033

aggatcttga ggctgggtgtt attcaggtta tccaaatcga cgaggctgca ctgagagaag 60  
 ggctgccgct ccgcaaggct gagcatgcat tctacttga ctgggctgtc cactccttca 120  
 gaattaccaa ctgcgagatc caggacacca ccagatcca gaccacatg tgctactcca 180  
 atttcaacga catcatccac tcgatcatcg acatggacgc ggacgtgatc accattgaga 240  
 actcggggtc cgacgagaag ctgctctccg tgttccgcga gggcgtcaag ta 292

<210> 1034  
 <211> 320  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1034

cttgacagcg agattaagtc ttggcttgct ttgctgccc agaaggttgt tgaggttgat 60  
 gctcttgcta aggcattggc tgggtcaaaag gatgaggctt acttcgcagc aaatgctgca 120  
 gctcaggcct cgaggaaatc ctcaccccggt gtgaccaatg aagaagtcca gaaggctgct 180  
 gctgctctca agggctctga ccaccgccgt gcgaccaacg ttagtgctag attggatgcc 240  
 cagcagaaga agcttaacct cccatcctt cccaccacta caatcggttc gttcccacaa 300  
 accgttgagc tcaggagggt 320

<210> 1035  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1035

acttcggtga gcagctctcc ggttttgcgt tcaactgccaa tggctgggtg cagtcttatg 60  
 ggtcaagggtg cgtcaagcct ccgatcatct acggtgacgt gagccgcccc aaccccatga 120  
 ccgttttctg gtcgaagacg gcacagagca tgacgtctcg cccaatgaag gggatgctga 180  
 ctggcccagt cacaatcctc aactggtcct tcgtccggaa tgaccagccg aggtttgaga 240  
 cttgctacca gatcgctctt gcgatcaaga aggaagtcga ggatcttgag gctgggtgga 300  
 ttcag 305

<210> 1036  
 <211> 278  
 <212> nucleic acid  
 <213> Zea mays

<400> 1036  
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 gagtacttcg gtgagcaatt atctggtttt gcgttcaactg ccaacggatg ggtgcaatcc 120  
 tatggatcac gctgtgtgaa gccacccatt atctatgggtg atgtcagccg gccgaacccc 180  
 atgactgttt tctgggtccaa gatggcacag agcatgaccc ctggtcccat gaagggaatg 240  
 ttgactggtc cggtcacaat cctgaactgg tcattcgt 278

<210> 1037  
 <211> 298  
 <212> nucleic acid  
 <213> Zea mays

<400> 1037  
 ctctgaggag gaatacgtca ctgccatcaa ggaggaaatc aacaaggctg ttaagcttca 60  
 agaagagctt gacattgatg tgcttggtgca tggcgagcct gagagaaacg atatggttga 120  
 gtacttcggt gagcagctct ccggttttgc gttcactgcc aatggctggg tgcagtctta 180  
 tgggtcaagg tgcgtcaagc ctccgatcat ctacggtgac gtgagccgcc ccaaccccat 240  
 gaccgttttc tggtcgaaga cggcacagag catgacgtct cgcccaatga aggggatg 298

<210> 1038  
 <211> 298  
 <212> nucleic acid  
 <213> Zea mays

<400> 1038

ctctgaggag gaatacgtca ctgccatcaa ggaggaaatc aacaaggctcg ttaagcttca 60  
agaagagctt gacattgatg tgcttgtgca tggcgagcct gagagaaaacg atatggttga 120  
gtacttcggt gaggagctct ccggttttgc gttcactgcc aatggctggg tgcagtctta 180  
tgggtcaagg tgcgtcaagc ctccgatcat ctacggtgac gtgagccgcc ccaaccccat 240  
gaccgttttc tggtcgaaga cggcacagag catgacgtct cgcccaatga aggggatg 298

<210> 1039

<211> 447

<212> nucleic acid

<213> Zea mays

<400> 1039

ggcgctcggg gttgataccg tgccagtact tgttggacca gtctcgtaac tgttgcngnn 60  
aaagcctgcc aagggtgtgg agaagggtatt cctcttctt tcccttctta gcagcatcct 120  
cccagtctac aaggagggtca ttgctgagtt gaaggcagct ggggcttcat ggattcagtt 180  
tgatgagccc actcttgtcc tcgaccttga ttctgacaaa ttggtgcat tctctgctgc 240  
atatgcagaa cttgaatctg tactttctgg attgaatgtg cttgttgaga cttactttgc 300  
tgatgttcct gctgagtcct acaagaccct aacatccctg aacagtgtga ctgcttatgg 360  
gtttgatent gtccggggac ccnaaactnt tgggcttggc accaaggctg gnttccttnt 420  
tggaanaaac cctntttgct ggggggtg 447

<210> 1040

<211> 301

<212> nucleic acid

<213> Zea mays

<400> 1040

ctccaagaag agcttgacat tgatgtgctt gtgcatggcg agcctgagag aaacgatatg 60  
gttgagtact tcggtgagca gctcctccgg ttttgcgttc actgccaatg gctgggtgca 120  
gtcttatggg tcaaggtgcg tcaagcctcc gatcatctac ggtgacgtga gccgccccaa 180  
ccccatgacc gttttctggt cgaagacggc gcagagcatg acgtctcgcc caatgaaggg 240  
gatgctgact ggcccagtca caatcctcaa ctggctcttc gtccggaatg accagccgag 300



<210> 1041  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays

<400> 1041

caaggtcggtt aagctccaag aagagcttga cattgatgtg cttgtgcatg gcgagcctga 60  
 gagaaacgat atggttgagt acttcggtga gcagctctcc ggttttgcgt tcaactgccaa 120  
 tggctgggtg cagtcttatg ggtcaagggtg cgtcaagcct ccgatcatct acggtgacgt 180  
 gagccgcccc aaccccatga ccgtttttctg gtccaagagg cgcagagcat gacgtctcgc 240  
 ccaatgaagg ggatgctgac tggcccagtc acaatcctca actggtcctt cgtccgg 297

<210> 1042  
 <211> 548  
 <212> nucleic acid  
 <213> Zea mays

<400> 1042

gnnnaggtgn tnannggggn nggaaggaac nttttaccgg tccggaattc ccgggtcgac 60  
 ccacgcgtcc gccacgcgt ccgctcgatc atcgacatgg acgcggacgt gatcaccatt 120  
 gagaactcgc ggnccgacga gaagctgctc tccgtgttcc gcgagggcgt caagtacggc 180  
 gcgggcatcg gccccggtgt ctacgacatc cactccccca ggatcccgtc ggctgaggag 240  
 atcgccgacc gcatcgacaa gatgctggcg gtgctggaca ccaacatcct ctgggtgaac 300  
 cccgactgcg gcctcaagac ccgcaagtac acggaggta agcccgctg accaacaatgg 360  
 tctccgccgc taagctcatc cgcacccagc ttgncagcgc caagtgaagca ccnttttttt 420  
 tgcccttttc ggtttcgaag aaggcgtcgt cgatgccaat ttggtttcca ataaaaaggc 480  
 tttgcccgcc gtctgtgnac tccgctgngg taaggtaaga agttttcttaa aaaaaaaaaa 540  
 angggggc 548

<210> 1043  
 <211> 303  
 <212> nucleic acid  
 <213> Zea mays

<400> 1043

gacggcgag agcatgacgt ctgcccgaat gaaggggatg ctgactggcc cagtcacaat 60  
cctcaactgg tccttcgtcc ggaatgacca gccgaggttt gagacttgct accagatcgc 120  
tcttgcatc aagaaggaag tcgaggatct cgaggctggt ggtattcagg ttatccaaat 180  
cgacgaggct gcactgagag aagggtgcc gctccgcaag gctgagcatg cattctactt 240  
ggactgggct gtccactcct tcagaatcac caactgcgag atccaggaca ccaccagat 300  
cca 303

<210> 1044

<211> 602

<212> nucleic acid

<213> Zea mays

<400> 1044

gcggacgtga tcaccattga gaactcgcgg tccgacgaga agctgctctc cgtgttcgcg 60  
gagggcgctca agtacggcgc gggcatcggc cccggtgtct acgacatcca ctccccaggg 120  
atcccgtcgg ctgaggagat cgccgaccgc atcgacaaga tgctggcggt gctggacacc 180  
aacatcctct ggggtgaacc cgactgcggc ctcaagacc gcaagtacac ggaggtcaag 240  
cccgccctga ccaacatggt ctccgccgct aagctcatcc gcaccagct cgccagcgcc 300  
aagtgagcac cttttttttg cttttttcgt ttccgaggag ggcgtcgtcg atgccaattt 360  
gtttccaata aacagggctg tgcccgccgt tctgttgtag tccgtctgtg gttagggttag 420  
tagttttctt gatctcgccc ccacgcggtg cctgttttac tactctctgt ttggtggttt 480  
ctgaggcaag ttgcccggtg acttgatcgc taaaaaccga gtgggacatc tgcacggtt 540  
catctgncac cactnccccg aggtgtgctg gttcacggct caaagaaaga tgtcntatcg 600  
gc 602

<210> 1045

<211> 273

<212> nucleic acid

<213> Zea mays

<400> 1045

ggttgtcaag atccaagagg agcttgacat tgatgtgctt gtgcatggag agccagagag 60

aaatgacatg gttgagtact tcggtgagca attatctggt tttgcgttca ctgccaacgg 120  
 atgggtgcaa tcctatggat cacgctgtgt gaagccaccc attatctatg gtgatgtcag 180  
 ccggccgaac cccatgactg ttttctggtc caagatggca cagagcatga cccctcgtcc 240  
 catgaaggga atgttgactg gtccggtcac aat 273

<210> 1046  
 <211> 328  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1046

atcatcgccg ctgcgcgctc tcctccgctc ccgtcgtccg gaaagatggc gtcccacatt 60  
 gttggatacc ctgcgatggg ccccaagagg gagctcaagt ttgcacttga gtctttctgg 120  
 gatgggaaga gcaccgctga ggatctggag aagggttgcta ccgacctcag ggccagtatc 180  
 tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaacacttt ctgtactat 240  
 gaccagggtc ttgacaccac tgccatgctc ggcgctgtcc ccgaacgtta ctcatgnact 300  
 ggaggagaga ttggatttga cacctact 328

<210> 1047  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1047

aaaaaagatc tctgaggagg aatacgtcac tgccatcaag gaggaaatca acaaggctgt 60  
 taagcttcaa gaagagcttg acattgatgt gcttgtgcat ggcgagcctg agagaaacga 120  
 tatggttgag tacttcggtg agcagctctc cggtttggcg ttactgcca atggctgggt 180  
 gcagtcttat gggtaaggt gcgtcaagcc tccgatcatc tacggtgacg tgagccgccc 240  
 caaccccatg accgttttct ggtcgaagac ggcacagagc atgacgtctc gcccaat 297

<210> 1048  
 <211> 264  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1048

ctcttttcaga atcaccaact ggggagtcca ggacaccacc cagatccaca cccatatgtg 60  
ctactccaac ttcaacgaca tcatccactc catcatcgac atggacgcag atgtgatcac 120  
catcgagaac tcccgggtccg acgagaagct actgtccgtg ttccgcgagg gtgtgaagta 180  
cggagctgga atcgggtccag gtgtctacga catccactcg cccaggatcc cctccaccga 240  
ggagatcgcg gaccgcgtca acaa 264

<210> 1049  
<211> 471  
<212> nucleic acid  
<213> Zea mays

<400> 1049

gggnnnnacg ttantcngcg ngnnccgtca naggacncac nggccgaggc ncgcgtccac 60  
accnatcatc gccgctgcgc cgtctcctcc gctcccgtcg tccggaaaga tggcgtccca 120  
cattgttggg taccctcgca tgggccccaa gagggagctc aagtttgcaac ttgagtcttt 180  
ctgggatggg aagagcaccg ctgaggatct ggagaagggt gctaccgacc tcagggccag 240  
tatctggaag cagatggctg atgctgggat caagtacatc cccagcaaca ccttctcgta 300  
ctatgaccag gtccttgaca ccactgccat gctcggcgct gtccccgaac gctactcatg 360  
gactggagga gagattgggt ttgacaccta acttctccat ggccaagggc aacgccactg 420  
tccctgctat gggagatacc aanttgggtt tggacacaaa ctaccaactt t 471

<210> 1050  
<211> 291  
<212> nucleic acid  
<213> Zea mays

<400> 1050

cgctcccgtc gtccggaaag atggcgctccc acattgttgg ataccctcgc atgggccccca 60  
agaggagct caagtttgca cttgagtctt tctgggatgg gaagagcacc gctgaggatc 120  
tggaagaagg tgctaccgac ctgaggcca gtatctggaa gcagatggct gatgctggga 180  
tcaagtacat cccagcaac actttctcgt actatgacca ggtccttgac accactgcca 240  
tgctcggcgc tgtccccgaa cgttactcat ggactggagg agagattgga t 291

<210> 1051

<211> 349  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1051  
  
 atccactcca tcatcgacat ggatgccgat gtgatcacga tcgagaactc ccgatctgac 60  
 gagaagctna ctgtccgtct tccgtgaggg tgtgaagtac ggagctgggc attggccctg 120  
 gtgtctacga catccactct cctaggattc acctccaccg aggagatcgc agaccgcgtc 180  
 gagaagatgc tcgccgtgct ggacaccaac atcctctggg tgaaccctga ctgtggtctc 240  
 aanacacgca agtacacgga ggtcaagccc gccctgacca acatggtctc ggccancaaag 300  
 ctcacccgca cccagcttgc caacgcgaaa tgagtcgttt gatagctcc 349

<210> 1052  
 <211> 333  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1052  
  
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 tctcgtacta tgaccaggct cttgatacca cggccatgct tgggtgctgtc ccagagcgct 120  
 actcttggtgac tggaggcgag attggcttga gcacctatct ctcaatggcc aggggaaatg 180  
 ccactgtccc tgctatggag atgaccaagt ggtttgatac aaactaccac tttattgtcc 240  
 ctgaattggc ccaagcacca agttcacata cgcttctcac aangctgttt ctgagtacaa 300  
 ggaggcaaaa gcgctcggtt ttgatatgtc cca 333

<210> 1053  
 <211> 299  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1053  
  
 ctggaaagta cctcttttgc ggtgttgtgg atggacgcaa catctgggct gatgatcttg 60  
 ctacatctct cagcactctc cagtctcttg aggtgttgtt tgggaaggac aagcttgctg 120  
 tatcaacttc ctgctcgctc atgcacaccg ctgtggatct tgtgaacgag actaagcttg 180  
 acagcgagat taagtcttgg cttgcttttg ctgcccagaa ggttgttgag gttgatgctc 240

ttgctaaggc attggctggt caaaaggatg aggcttactt cgcagcaa at gctgcagct 299

<210> 1054  
 <211> 281  
 <212> nucleic acid  
 <213> Zea mays

<400> 1054

gctaccagat cgctcttgcg atcaagaagg aagtcgagga tcttgaggct ggtggtattc 60  
 aggttatcca aatcgacgag gctgcactga gagaagggct gccgctccgc aaggctgagc 120  
 atgcattcta cttggactgg gctgtccact cttcagaat taccaactgc gagatccagg 180  
 acaccacca gatccacacc cacatgtgct actccaactt caacgacatc atccactcga 240  
 tcatcgacat ggacgoggac gtgatcacca ttgagaatcg g 281

1054  
 281  
 nucleic acid  
 Zea mays  
 1055  
 305  
 nucleic acid  
 Zea mays  
 1055  
 60  
 120  
 180  
 240  
 300  
 305

<210> 1055  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays

<400> 1055

cgccgctgcg ccgtctctc cgtctccgctc gtccggaaag atggcgtccc acattgttgg 60  
 ataccctcgc atggggccca agagggagct caagtttgca cttgagtctt tctgggatgg 120  
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180  
 gcagatggct gatgctggga tcaagtacat cccagcaac actttctcgt actatgacca 240  
 ggtccttgac accactgcc tgcctggcgc tgtccccgaa cgttactcat ggactggagg 300  
 agaga 305

<210> 1056  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays

<400> 1056

ggctggtcaa aaggatgagg ctacttctgc agcaa atgct gctgctcagg cctcgaggaa 60  
 atcctcacc cgtgtgacca atgaagaagt ccagaaggct gctgctgctc tcaagggctc 120  
 tgaccaccgc cgtgggacca acgttagtgc tagattggat gccagcaga agaagcttaa 180

cctccccatc cttcccaacca ctacaatcgg ttcgttccca caaaccgttg agtcaggag 240  
 ggtccgccgt gagtacaagg caaaaaagat ctctgaggag gaatacgtca ct 292

<210> 1057  
 <211> 279  
 <212> nucleic acid  
 <213> Zea mays

<400> 1057

atcaagaagg aagtcgagga tcttgaggct ggtggtattc aggttatcca aatcgacgag 60  
 gctgcaactga gagaagggct gccgctccgc aaggctgagc atgcattcta cttggactgg 120  
 gctgtccact ccttcagaat taccaactgc gagatccagg acaccacca gatccacacc 180  
 cacatgtgct actccaactt caacgacatc atccactcga tcatcgacat ggacgcggac 240  
 gtgatcacca ttgagaactc gcggtccgac gagaagctg 279

<210> 1058  
 <211> 286  
 <212> nucleic acid  
 <213> Zea mays

<400> 1058

atcgtctcttg cgatcaagaa ggaagtcgag gatctcgagg ctggtggtat tcaggttatc 60  
 caaatcgacg aggctgcaact gagagaaggg ctgccgctcc gcaaggctga gcatgcattc 120  
 tacttggact gggctgtcca ctcttcaga atcaccaact gcgagatcca ggacaccacc 180  
 cagatccaca cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgac 240  
 atggacgcgg acgtgatcac catcgagaac tcgcggtccg acgaga 286

<210> 1059  
 <211> 269  
 <212> nucleic acid  
 <213> Zea mays

<400> 1059

caagatccaa gaggagcttg acattgatgt gcttgtgcat ggagagccag agagaaatga 60  
 catggttgag tacttcggtg agcaattatc tggttttgcg ttactgcca acggatgggt 120  
 gcaatcctat ggatcacgct gtgtgaagcc acccattatc tatggtgatg tcagccggnc 180

gaaccccatg actgttttct ggtccaagat ggcacagagc atgacccctc gtcccatgaa 240  
 gggaatgttg actggtccgg tcacaatcc 269

<210> 1060  
 <211> 323  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1060

gcaacatctg gggctgatga tcttgctaca tctctcagca ctctccagtc tcttgaggct 60  
 gttgttgga aggacaagct tgtcgtatca acttcctgct cgctcatgca caccgctgtg 120  
 gatcttgta acgagactaa gcttgacagc gagattaagt cttggcttgc ttttgctgcc 180  
 cagaagggtg ttgagggtga tgctcttgct aaggcattgg ctggtcaaaa ggatgaggct 240  
 tacttcgcag caaatgctgc agctcaggcc tcgaggaaat cctcaccocg tgtgaccaat 300  
 gaagaagtcc agaaggctgc tgc 323

<210> 1061  
 <211> 287  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1061

angatcttga ggctgctggt attcaggtga tccagattga tgaggcagct ctaaggagg 60  
 gtctgccact gcgcaagtta gagcatgcat tctacctgga ctgggctgtc cactctttca 120  
 gaatcaccaa ctgcggagtc caggacacca cccagatcca caccatattg tgctactcca 180  
 attcaanga catcatccac tccatcatcg acatggacgc agatgtgatc accatcgaga 240  
 actcccgtc cgacgagaag ctactgtccg tgttccgcga ggtgtga 287

<210> 1062  
 <211> 295  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1062

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 agcaccgctg aggatntgga gaagggtgct accgacctca gggccagtat ctggaagcag 120



atggctgatg ctgggatcaa gtacatcccc agcaacacct tctcgtacta tgaccaggtc 180  
 cttgacacca ctgccatgct cggcgctgtc cccgaacgct actcatggac tggaggagag 240  
 attgggtttg acacctactt ctccatggcc aggggcaacg ccactgtccc tgcta 295

<210> 1063  
 <211> 304  
 <212> nucleic acid  
 <213> Zea mays

<400> 1063

gtcagagcat gcattotacc tggactgggc tgtccactct ttcaggatca ccaactgcgg 60  
 agtccaggac accacccaga tccacacca catgtgctac tccaacttca acgacatcat 120  
 ccaactccatc atcgacatgg atgccgatgt gatcacgac gagaaactccc gatctgacga 180  
 gaagctactg tccgtotacc gtgagggtgt gaagtacgga gctggcattg gccctgggtg 240  
 ctacgacatc catctcttag gattccctcc accgaggaga tcgcagaccg cgtcgagaag 300  
 atgc 304

<210> 1064  
 <211> 293  
 <212> nucleic acid  
 <213> Zea mays

<400> 1064

ctgcgagatc caggacacca cccagatcca caccacatg tgctactcca acttcaacga 60  
 catcatccac tcgatcatcg acatggacgc ggacgtgatc accatcgaga actcgcggtc 120  
 cgacgagaag ctgtctctcc tgttccgggg gtgtcaagta cggcgcgggc atcggccccg 180  
 gtgtctacga catccactcc cccaggatcc cgtcggccga ggagatcgcc gaccgcatcg 240  
 acaagatgct ggccgtgctg gacaccaaca tcctctgggt gaaccccgac tgc 293

<210> 1065  
 <211> 285  
 <212> nucleic acid  
 <213> Zea mays

<400> 1065

tcttggttg cttttgctgc ccagaaggtt gttgaggttg atgtctctgc taaggcattg 60

gctgggtcaaa aggatgagggc ttacttcgca gcaaagtctg ctgctcaggc ctcgaggaaa 120  
tcctcacccc gtgtgaccaa tgaagaagtc cagaaggctg ctgctgctct caagggctct 180  
gaccaccgcc gtgggaccaa cgttagtgtg agattggatg cccagcagaa gaagcttaac 240  
ctcccatcc ttcccaccac tacaatcggt tcgttcccac aaacc 285

<210> 1066  
<211> 285  
<212> nucleic acid  
<213> Zea mays  
<400> 1066

gggctgtcca ctcttcaga attaccaact gcgagatcca ggacaccacc cagatccaca 60  
cccatatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgac atggacgcgg 120  
acgtgatcac cattgagaac tcgcgggtccg acgagaagct gctctccgtg ttccgcgagg 180  
gcgtcaagta tggcgcgggc atcggtcccg gtgtctacga catccactcc cccaggatcc 240  
cgtcggctga ggagatcgcc gaccgcatcg acaagatntg gcggt 285

<210> 1067  
<211> 277  
<212> nucleic acid  
<213> Zea mays  
<400> 1067

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atgaccaggt ccttgacacc actgccatgc tcggcgctgt cccgaacgt tactcatgga 120  
ctggaggaga gattggattt gacacctact tctccatggc caggggcaac gccactgttc 180  
ctgctatgga gatgaccaag tggtttgaca ccaactacca ctttattgtc cctgaattgg 240  
gcccaaacac caagttctcc tatgcttctc acaaggc 277

<210> 1068  
<211> 280  
<212> nucleic acid  
<213> Zea mays  
<400> 1068

cccagatcca caccacatg ngcncgtcca acttcaacga catcatccac togatcatcg 60

acanggacgc ggacgtgac accatcgaga actcgcggtc cgacgagaag ctgctctccg 120  
 tgttccgtga ggggtgtcaag tacggcgcggtc gcatcggccc cgggtgtctac gacatccact 180  
 cccccaggat cccgtcggcc gaggagatcg ccgaccgcat cgacaagatg ctggccgtgc 240  
 tggacaccaa catcctctgg gtgaaccccg actgcggcct 280

<210> 1069  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1069

cattgatntg cttgtgcatg gcgagcctga nagaaacgat atggttgagt acttcggtga 60  
 gcagctctcc ggttttgcgt tcaactgcaa tggctgggtg cagtcttatg ggtcaagggtg 120  
 cgtcaagcct ccgatcatct acggtgacgt gagccgcccc aaccccatga ccgttttctg 180  
 gtcgaagagg cgcagagcat gacgtctcgc ccaatgaagg ggatgctgac tggncagtc 240  
 acaatcctca actgggtcctt cgtccggaat gaccagccga ggtttgaaac tgcnaac 297

<210> 1070  
 <211> 289  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1070

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 tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 120  
 gctctccggt tttgcgttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 180  
 caagcctccg atcatctacg gtgacgtgag ccgccccaac cccatgaccg ttttctggtc 240  
 gaagaggcac agagcatgac gtctcgccca atgaaggga tgctgactg 289

<210> 1071  
 <211> 486  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1071

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gaaacggtgg aactcaggng ggttcgncgt gaatacaagg caaagaagat caccgaggac 120  
gaatacatca gtgccatcaa ggaagaaatc agcaaggtn gcaagatcca ataggagctn 180  
gacattgatg tgcttgtgca tggagagcca gagagaaatg acatgggtga gtacttcggn 240  
gagcaattat ctgggttttag cgttcactgc caacggatgg gtgcaatnct attgatcacg 300  
ctgtggtgaa gccacccatt atctatgnga tgtcaaagcg gcccnacccc atnctgntnt 360  
ctgggtccaa gatngancat acatgacccc tttnccatg naagggaatg gttgactggg 420  
tcnggtcaca attcttaaac ttggtcattn gntaaaggaa cgacnancct taggtttnaa 480  
anannc 486

<210> 1072  
<211> 286  
<212> nucleic acid  
<213> Zea mays

<400> 1072  
aaccocatga ccgttttctg gtcgaagagg tcacagagca tgacgtctcg cccaatgaag 60  
gggatgctga ctggcccagt cacaatcctc aactggctct tcgtccggaa tgaccagccg 120  
aggtttgaga cttgctacca gntcgctctt gcgatcaaga aggaagtcca ggatcttgag 180  
gctggtggta ttacaggttat ccaaatcgac gaggtgcac tgagagaagg gctgccgctc 240  
cgcaangctg agcatgcatt ctacttgac tgggctgtcc actcct 286

<210> 1073  
<211> 305  
<212> nucleic acid  
<213> Zea mays

<400> 1073  
gaagagcttg acattgatgt gcttgtgcat ggcgagcctg agagaaacga tatggttgag 60  
tacttcggtg agcagctctc cggttttgctg ttactgcca atggctgggt gcagtcttat 120  
gggtcaaggt gcgtcaagcc tccgatcatc tacggtgacg tgagccgccc caaccccatg 180  
accgttttct ggtcgaagac ggcacagagc atgacgtctc gcccaatgaa ggggatgctg 240  
actggcccag tcacaatcct caactggtct tcgtccggaa tgaccagccg aggtttgaga 300  
ttgct 305

<210>	1074
<211>	291
<212>	nucleic acid
<213>	Zea mays

caatcgggttc	gttcccacaa	accgttgagc	tcaggagggt	cgcgcngag	tacaaggcaa	60
aaaagatctc	tgaggaggaa	tacgtcactg	ccatcaagga	ggaaatcaac	aaggtcgtta	120
agcttcaaga	agagcttgac	attgatgtgc	ttgtgcatgg	cgagcctgag	agaaacgata	180
tggttgagta	cttcggtgag	cagctctccg	gttttgcggt	cactgccaat	ggctgggtgc	240
agtottatqg	gtcaaggngc	qncaagcctc	cgatcatnga	gggtgaccnc	c	291

<211>	297
<212>	nucleic acid
<213>	Zea mays

tgcacttgag	nctttctggg	atgggaagag	cacgcgtgag	gatctggaga	aggttgctac	60
cgacctcagg	gccagtatct	ggaagcagat	ggctgangct	gggatcaagt	acatccccag	120
caacactttc	tcgtactatg	accaggtcct	tgacaccact	gccatgctcg	gcgctgtccc	180
cgaacgttac	tcatggactg	gaggagagat	tggatttgac	acctacttct	ccnnggccag	240
gggcaacgcc	actgttctctg	ctatggagat	gaccangtgg	tttgacacca	ataccac	297

<211>	312
<212>	nucleic acid
<213>	Zea mays

acctatcattc	gccgctgccc	cgtctcctcc	gctcccgtcg	tccggaaaaga	tggcgtccca	60
cattgttgga	taccctcgca	tgggccccaa	gaggggagctc	aagtttgcac	ttgagctttt	120
ctgggatggg	aagagcaccg	ctgaggatct	ggagaagggt	gctaccgacc	tcagggccag	180
tatctggaag	cagatggctg	atgctgggat	caagtacatc	cccagcaaca	ccttctcgta	240
ctatgaccag	gtccttgaca	ccactgccat	gctcggcgct	gtccccgaac	gctactcatg	300

gactggagga ga

312

<210> 1077  
<211> 307  
<212> nucleic acid  
<213> Zea mays

<400> 1077

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cttgtgaacg agactaagct tgacagcgag attaagtctt ggcttgcctt tgctgcccag 120  
aaggttgttg aggggtganc tcttgctaag gcantggctg gtcaaaagga tgaggcttac 180  
ttcgcagcaa atgctgctgc tcaggcctcg aggaaatcct caccocgtgt gaccaatgaa 240  
gaagtccaga aggctgctgc tgctctcaag ggctctgacc accgccgtgg gaccaacgtt 300  
agtgcta 307

<210> 1078  
<211> 274  
<212> nucleic acid  
<213> Zea mays

<400> 1078

ggcgtccac attgttggat accctcgcat gggccccaag anggagctca agtttgcact 60  
tgagtctttc tgggatggga agaacaccgc tgagnatctg gagaagggtg ctaccgacct 120  
cagggccagt atctggaagc agatggctga tgctgggatc aagtacatcc ccagcaacac 180  
tttctogtac tatgaccagg tcttgacac cactgccatg ctacgcgtg tccccgaacg 240  
ttactcatgg actggaggag agattggatt tgac 274

<210> 1079  
<211> 289  
<212> nucleic acid  
<213> Zea mays

<400> 1079

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tggagatgac caagtgggtt gacaccaact accactttat tgtccctgaa ttgggcccac 120  
acaccaagtt ctctatgct tctcacaagg ctgttaacga atataaggag gctaaggcgc 180

tcggtgttga taccgtgcc a gtacttgttg gaccagtctc gtacctgttg ctctcaaagc 240  
 ctgccaaagg tgtggagaag ggattccctc ttctttccct tcttagcag 289

<210> 1080  
 <211> 286  
 <212> nucleic acid  
 <213> Zea mays

<400> 1080

ctaagcttga cagcgagatt aagtcttggc ttgcttttgc tgcccagaag gttgttgagg 60  
 ttgatgctct tgctaaggca ttggctggtc aaaaggatga ggcttacttc gcagcaaattg 120  
 ctgcagctca ggccctcgagg aaatcctcac cccgtgtgac caatgaagaa gtccagaagg 180  
 ctgctgctgc tctcaagggc tctgaccacc gccgtgagac caacgttagt gctagattgg 240  
 atgccacgca gaagaagctt aacctcccca tccttccac cactac 286

<210> 1081  
 <211> 323  
 <212> nucleic acid  
 <213> Zea mays

<400> 1081

cgccgctgag ccgtctctc cgctcccgtc gtccggaaag atggcgctcc acattgttgg 60  
 ataccctcgc atggggccca agagggagct caagtttgca cttgagtctt tctgggatgg 120  
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180  
 gcagatggct gatgctggga tcaagtacat cccagcaac accttctcgt actatgacca 240  
 ggtccttgac accactgcc a ngctcggcgc tgtccccgaa cgntactcat ggactgggng 300  
 gagaaattgg gtttcacacc tat 323

<210> 1082  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays

<400> 1082

cacaaaccgt tgcagctcag gaggggtccgc cgtgagtaca aggcaaaaaa gatctctgag 60  
 gaggaatacg tcaactgcat caaggaggaa atcaacaagg tcgttaagct ccaagaagag 120

cttgacattg atgtgcttgt gcatggcgag cctgagagaa acgatatggg tgagtacttc 180  
 ggtgagcagc tctccggttt tgcgttcact gccaatggct ggggtgcagtc ttatgggtca 240  
 aggtgcgtca agcctccgat catctacggg gactgagccg cccaacccc atgaccgttt 300  
 tctgg 305

<210> 1083  
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 <212> nucleic acid  
 <213> Zea mays  
 <400> 1083

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 ganaaacgat atggttgagt acttcgggtga gcagctctcc ggttttgctg tcaactgcca 180  
 tggttgggtg cagtcttatg ggtcaagggt cgtcaagcct ccgatcatct acggtgacgt 240  
 gagccgcccc aaccccatga ccgttttctg gtcgnagacg g 281

<210> 1084  
 <211> 314  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1084

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 gatctcgagg ctggtggtat tcaggttatc caaatcgacg aggctgcact gagagaaggg 120  
 ctgcgcctcc gcaaggctga gcatgcattc tacttggacn gggctgtcca cnccttcaga 180  
 atcaccaacn gcgagatcca ggacaccacc cagatccaca cccacatgtg ctacnccaac 240  
 ttcaacgaca tcatccactc gatcatcgac atggacgcgg ngtgatcacc acgaganncn 300  
 ggtcgcgaga actg 314

<210> 1085  
 <211> 281  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1085



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 gtgggaccaa cgttagtgt agattggatg cccagcagaa gaagcttaac ctccccatcc 120  
 ttccccaccac tacaatcggg tggttcccac aaaccgttga gtcaggagg gtccgccgtg 180  
 agtacaaggc aaaaaagatc tctgaggagg aatacgtcac tgccatcaag gaggaaatca 240  
 acaaggctgt taagcttcaa gaagagcttg acattgatgt g 281

<210> 1086  
 <211> 306  
 <212> nucleic acid  
 <213> Zea mays

<400> 1086

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 ttgttgata ccctcgcatg ggccccaaga gggagctcaa gtttgcaatt gagtctttct 120  
 gggatgggaa gagcacccgt gaggatctgg agaaggttgc taccgacctc agggccagta 180  
 tctggaagca gatggctgat gctgggatca agtacatccc cagcaacacc ttctcgtact 240  
 atgaccaggt ccttgacacc actgccatgc tcggcgctgt ccccgaaacgc tactcatgga 300  
 ctggag 306

<210> 1087  
 <211> 265  
 <212> nucleic acid  
 <213> Zea mays

<400> 1087

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 cggagtccag gacaccaccc agatccacac ccacatgtgc tactccaact tcaacgacat 180  
 catccactcc atcatcgaca tggatgccga tgtgatcacg atcgagaact cccggtctga 240  
 cgagaagcta ctgtccgtct tccgt 265

<210> 1088  
 <211> 266  
 <212> nucleic acid  
 <213> Zea mays

<400> 1088

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caggagggtc cgccgtgagt acaaggcaaa aaagatctct gaggaggaat acgtcactgc 120

catcaaggag gaaatcaaca aggtcgttaa gcttcaagaa gagcttgaca ttgatgtgct 180

tgtgcatggc gagcctgaga gaaacgatat ggttgagtac ttcggtgagc agctctccgg 240

ttttgcgttc actgccaatg gctggg 266

<210> 1089

<211> 311

<212> nucleic acid

<213> Zea mays

<400> 1089

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gcgtcaagcc tccgatcctc tacgggtgacg tgagccgccc caaccccatg accgttttct 120

ggtcgaagac ggacacagagc atgacgtctc gcccaatgaa ggggatgctg actggcccag 180

tcacaatcct caaotggctc ttcgtccgga atgaccagcc gaggtttgag acttgctacc 240

agatcgctct tgcgatcaag aaggaagtcg aggatttgag gctggtggta ttcaggttat 300

ccaaatcgac g 311

<210> 1090

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 1090

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ggatgggaag agcaccgctg aggatctgga gaaggttgct accgacctca gggccagtat 180

ctggaagcag atggctgatg ctgggatcaa gtacatcccc agcaacacct tctcgtacta 240

tgaccaggtc cttgacacca ctgccatgct cggcgtgtc cccgaacgct antcatggac 300

tgaggaga 309

<210> 1091

<211> 291  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1091  
  
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 tgctctccgt gttccgtgag ggtgtcaagt acggcgcggg catcgggccc ggtgtctacg 120  
 acatccactc cccaggatc ccgtcggccg aggagatcgc cgaccgcatc gacaagatgc 180  
 tggccgtgct ggacaccaac atcctctggg tgaacccga ctgcggcctc aagacccgca 240  
 agtacacgga ggtcaagccc gccctgacca acatggtctc cgctgctaag t 291

<210> 1092  
 <211> 295  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1092  
  
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 tggcgagcct gagagaaaacg atatggttga gtacttcggt gagcagctct ccggttttgc 180  
 gttcactgcc aatggctggg tgcagtctta tgggtcaagg tgcgtcaagc ctccgatcat 240  
 ctacggtgac gtgagccgcc ccaaccccat gaccgttttc tggtcgaaga ggcac 295

<210> 1093  
 <211> 285  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1093  
  
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 agtcacaatc ctcaactggt ccttcgtccg gaatgaccag ccgaggtttg agacttgcta 180  
 ccagatcgct cttgcgatca agaaggaagt cgaggatctc gaggctggtg gtattcaggt 240  
 tatccaaatc gacgaggctg cactgagaga agggctgccg ctccg 285

<210> 1094

<211> 266  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1094  
  
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 cacatgtgct actccaactt caacgacatc atccactcga tcatcgacat ggacgaggac 180  
 gtgatcacca ttgagaactc gcggtccgac ganaagctgc tctccgtgtt ccgcgagggc 240  
 gtcaagtacg gcgcggggcat cggccc 266

<210> 1095  
 <211> 327  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1095  
  
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 tgagacttgc taccagatcg ctcttgcat caagaaggaa gtcgaggatc ttgaggctgg 120  
 tggatttcag gttatccaaa tcgacgaggc tgcactgaga gaagggtgc cgctccgcaa 180  
 ggctgagcat gcattctact tggactgggc tgtccactcc ttcagaatta ccaactgoga 240  
 gatccaggac accacccaga tccacacca catgtgctat ccaattcacg acatcatcca 300  
 togatcatcg ncatggngcg gcgtgat 327

<210> 1096  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1096  
  
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 ccatcatcga catggatgcc gatgtgatca cgatcgagaa ctcccgtct gacgagaagc 120  
 tactgtccgt cttccgtgag ggtgtgaagt acggagctgg cattggccct ggtgtctacg 180  
 acatccactc tcttaggatt ccttcaccg aggagatcg agaccgcgtc gagaagatgc 240  
 tcgngtgtn gananaant 259

<210> 1097  
 <211> 280  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1097  
  
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 aggtgtctac gacatccact cgcccaggat cccctccacc gaggagatcg cggaccgcgt 120  
 caacaagatg ctgcgcgtgc tcgacaccaa catcctctgg gtgaaccctg actgcggtct 180  
 caagacacgc aagtacacgg aggtgaagcc tgccctgacc aacatggtct ccgccaccaa 240  
 gtcacatcgc acccagctcg gcagcgcgaa atgagtcaga 280

<210> 1098  
 <211> 303  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1098  
  
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 atggcgctcc acattgttgg ataccctcgc atgggcccc aagaggagct caagtttgca 120  
 cttgagtctt tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac 180  
 ctcagggcc a gtatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac 240  
 actttctcgt actatgacca ggtccttgac accactgcc a tgctcggcgc tgtccccgaa 300  
 cgt 303

<210> 1099  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1099  
  
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 cccacattgt tggataccct cgcattgggc ccaagaggga gctcaagttt gcacttgagt 120  
 ctttctggga tgggaagagc accgctgagg atctggagaa ggttgctacc gacctcagg 180  
 ccagtatctg gaagcagatg gctgatgctg ggatcaagta catccccagc aacactttct 240

cgtactatga ccaggtcctt gacaccactg ccatgctcgg cgctgtcccc gaacggttact 300  
catgg 305

<210> 1100  
<211> 303  
<212> nucleic acid  
<213> Zea mays

<400> 1100

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cacattgttg gantaccctc gcatggggccc caagagggag ctcaagtttg cacttgagtc 120  
tttctgggat gggaagagca ccgtgagga tctggagaag gttgctaccg acctcagggc 180  
cagtatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca acactttctc 240  
gtactatgac caggggcaac gccactgttc ctgctatgga gatgaccaag tggtttgaca 300  
cca 303

<210> 1101  
<211> 288  
<212> nucleic acid  
<213> Zea mays

<400> 1101

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ttctggtcga agaggcacag agcatgacgt ctgcgccaat gaaggggatg ctgactggcc 120  
cagtcacaat cctcaactgg tcttcgtcc ggaatgacca gccgaggttt gagacttgct 180  
accagatcgc tcttgcatc aagaaggaag tcgaggatct tgaggctggt ggtattcagg 240  
ttatocaaat cgacgaggct gcactgagag aagggctgcc gctccgca 288

<210> 1102  
<211> 275  
<212> nucleic acid  
<213> Zea mays

<400> 1102

cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgnc catggacgcg 60  
gacgtgatca ccatcgagaa ctgcggtcc gacgagaagc tgctctccgt gttccgtgag 120

gggtgtcaagt acgggcgcggg catcgggcccc ggtgtctacg acatccactc cccagggatc 180  
 ccgtcgggcg aggagatcgc cgaccgcacg gacaagatgc tggccgtgct ggacaccaac 240  
 atcctctggg tgaaccccgga ctgcggcctc aagac 275

<210> 1103  
 <211> 319  
 <212> nucleic acid  
 <213> Zea mays

<400> 1103

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 tcgcccgaatg aaggggatgc tgactggccc agtcacaatc ctcaactggc ctttcgtccg 120  
 gaacgaccag ccgagggttg agacttgcta ccagatcgct cttgcgatca agaaggaagt 180  
 cgaggatctt gaggttggtg gtattcaggt tatccaaatc gacgaggctg cactgagaga 240  
 agggctgccc ctccgcaagg ctgagcatgc attctacttg gatgggctgt ccatccttca 300  
 gaataccaac tgcgagatc 319

<210> 1104  
 <211> 277  
 <212> nucleic acid  
 <213> Zea mays

<400> 1104

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 ctcaggcctc gaggaaatcc tcaccccgctg tgaccaatga agaagtccag aaggctgctg 120  
 ctgctctcaa gggctctgac caccgcccgtg ggaccaacgt tagtgctaga ttggatgccc 180  
 agcagaagaa gcttaacctc cccatccttc ccaccactac aatcggttcg tccccacaaa 240  
 ccgttgagct caggaggggtc cgccgtgagt acaangc 277

<210> 1105  
 <211> 308  
 <212> nucleic acid  
 <213> Zea mays

<400> 1105

caagaagagc ttgacattga tgtgcttgct catggcgagc ctgagagaaa cgatatgggtt 60

gagtacttcg gtgagcagct ctccggtttt gcgttcactg ccaatggctg ggtgcagtct 120  
tatgggtcaa ggtgcgtcaa gctccgcatc atctacgggtg angcnnaccg ccccaacccc 180  
atgaccgttt tctggctgaa gacggcacag agcatgacgt ctgcaccaat gaaggggatg 240  
ctgactggcc cagtcacaat cctcaatggc ccttcgtccg gaatgaccan ccgangtttg 300  
agattnct 308

<210> 1106  
<211> 313  
<212> nucleic acid  
<213> Zea mays

<400> 1106

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gttgctctca aagcccgcca aggggtgtga gaagggattc cctcttcttt cccttcttag 120  
cagcatcctc ccagtctaca aggaggtcat tgctgagttg aaggcagctg gggcttcatg 180  
gattcagttt gatgagccca ctcttgtcct cgaccttgat tctgacaaat tggttgcatt 240  
ctctgtgca taogcagaac ttgaatctgt actttctgga ttgaatgtgc ttgttgagac 300  
ttactttgct gat 313

<210> 1107  
<211> 279  
<212> nucleic acid  
<213> Zea mays

<400> 1107

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gtgaccaatg aagaagtcca gaaggctgct gctgctctca agggctctga ccaccgccgt 120  
gcgaccaacg ttagtgctag attggatgcc cagcagaaga agcttaacct ccccatcctt 180  
cccaccacta caatcgggtc gttcccacaa accgttgagc tcaggagggt ccgccgtgag 240  
tacaaggcaa aaaagatctc tgaggaggaa tacgtcact 279

<210> 1108  
<211> 264  
<212> nucleic acid  
<213> Zea mays



<400> 1108

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caactgttgct gctagattgg atgctcagca gaaaaagctc aaccttcctg tccttccac 120  
aaccacaatt ggttcattcc ctgagactgt ggaactcagg aggggtccgtc gtgagtacaa 180  
ggcaaagaag atctccgagg aggaatacat cagtgccatc aaggaagaaa tcagcaagggt 240  
tgtcaagatc caagaggagc ttga 264

<210> 1109

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 1109

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tgctgctgct caggcctcga ggaaatcctc acccctgtg accaatgaag aagtccagaa 120  
ggctgctgct gctctcaagg gctctgacca ccgccgtgg accaacgtta gtgctagatt 180  
ggatgccag cagaagaagc ttaacctccc catccttccc accactacaa toggttcgtt 240  
cccacaaacc gttgagctca ggagggtccg ccgtg 275

<210> 1110

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 1110

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cactgttctt gctatggaga tgaccaagtg gtttgacacc aactaccact ttattgtccc 120  
tgaattgggc ccaaacacca agttctccta tgcttctcac aaggctgtta acgaatataa 180  
ggaggctaag gcgctcgggtg ttgataccgt gccagtactt gttggaccag tctcgtacct 240  
gttgctctca aagcctgcc aaggtgtgga gaagggat 278

<210> 1111

<211> 310

<212> nucleic acid

<213> Zea mays

<400> 1111  
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 acattgtttg ataccctcgc atgggccccca agagggagct caagtttgca cttgagtctt 120  
 tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca 180  
 gtatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac accttctcgt 240  
 actatgacca ggtccttgac accactgcca tgctcggcgc tgtccccana acgtactca 300  
 tggactggag 310

<210> 1112  
 <211> 265  
 <212> nucleic acid  
 <213> Zea mays

<400> 1112  
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 gatgtgcttg tgcatggcga gcctgagaga aacgatatgg ttgagtactt cggtgagcag 120  
 ctctccggtt ttgcgttcac tgccaatggc tgggtgcagt cttatgggtc aagggtgcgtc 180  
 aagcctccga tcatctacgg tgacgtgagc cgccccaacc ccatgaccgt tttctgggtc 240  
 aagacggcgc agagcatgac gtctc 265

<210> 1113  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays

<400> 1113  
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 aggtgtctac gacatccact tcgcccagga tcccctccac cgaggagatc gcggaccgcg 120  
 tcaacaagat gctcgccgtg ctcgacacca acatcctctg ggtgaaccct gactgcggtc 180  
 tcaagacacg caagtacacg gaggtgaagc ctgccctgac caacatggtc tccgccacca 240  
 agctcatccg caccagctc ggcagcgcga aatgagtcag atttggtaga tccatggttt 300  
 gttga 305

<210> 1114

<211> 310  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1114  
  
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 ctatgcttct cacaaggctg ttaacgaata taaggaggct aaggcgctcg gtgttgatac 120  
 cgtgccagta cttgttggac cagtctcgta cctgttgctc tcaaagcctg ccaaggggtg 180  
 ggagaaggga ttccctcttc ttcccttct tagcagnact tcccagtcta caaggaggtc 240  
 attgctgagt tgaaggcagc tggggcttca ggattcagtt tgatgagccc atcttgtcct 300  
 cgactngatt 310

<210> 1115  
 <211> 331  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1115  
  
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 tatggttgag tacttcgggtg agcagctctc cggttttgcg ttcaactgcca atggcnggg 120  
 gcagtcttat ggggtcaagggt gcggtcaagcc tccgatcatc tacggtgacg tgagccgccc 180  
 caaccccatg accgttttct ggtcgaagac gggcacagag catgacgtct cgcaccaatga 240  
 angggatgct gactggccca gtcacaatcc tcnactggtc ttcggtccgga atgaccagcc 300  
 gangttgaga ttgctacnag atcgctcttg c 331

<210> 1116  
 <211> 279  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1116  
  
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 gagagaaacg atatggttga gtacttcgggt gagcagctct ccggttttgc gttcactgcc 120  
 aatggctggg tgcagtctta tgggtcaagg tgcgtcaagc ctccgatcat ctacggtgac 180  
 gtgagccgcc ccaaccccat gaccgttttc tggtcgaaga ggcacagagc atgacgtctc 240

gcccaatgaa ggggatgctg actggcccag tcacaatcc 279

<210> 1117  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays

<400> 1117

atcgccgctg cgccgtctcc tccgctcccg tcgtccggaa agatggcgtc ccacattggt 60  
 ggataccctc gcatggggcc ccaagaggga gctcaagttt gcacttgagt ctttctggga 120  
 tgggaagagc accgctgagg atctggagaa ggttgctacc gacctcaggg ccagtatctg 180  
 gaagcagatg gctgatgctg ggatcaagta catccccagc aacactttct cgtactatga 240  
 ccaggtcctt gacaccactg ccagtctcgg cgctgtcccc gaacgttact ca 292

<210> 1118  
 <211> 270  
 <212> nucleic acid  
 <213> Zea mays

<400> 1118

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 ggcgagcctg agagaaacga tatgggtgag tacttcggtg agcagctctc cggttttgcg 120  
 ttactgcca atggctgggt gcagtcttat gggtaaggt gcgtcaagcc tccgatcatc 180  
 tacggtgacg tgagccgccc caaccccatg accgttttct ggtcgaagag gcgcagagca 240  
 tgacgtctcg cccaatgaag gggatgctga 270

<210> 1119  
 <211> 252  
 <212> nucleic acid  
 <213> Zea mays

<400> 1119

cagcagaaga agcttaacct acccatcctt cncgccacta caatagggtc gttcccacaa 60  
 accgttgagc tcaggagggt ccgccgtgag tacaaggcaa aaaagatctc tgaggaggaa 120  
 tacgtcactg ccatcaagga ggaaatcaac aaggctgcta agtccaaga agagcttgac 180  
 attgatgtgc ttgtgcatgg cgagcctgag agaaacgata tggttgagta cttcggtgag 240

cagctctccg gt

252

<210> 1120  
<211> 284  
<212> nucleic acid  
<213> Zea mays

<400> 1120

cgttttcttg tcgaagaggc acagagcatg acgtctcgcc caatgaaggg gatgctgact 60  
ggcccagtc caatcctcaa ctggtccttc gtccggaatg accagccgag gtttgagact 120  
tgctaccaga togtctcttg gatcaagaag gaagtcgagg atcttgaggc tggtggtatt 180  
caggttatcc aaatcgacga ggctgcactg agagaagggc tgccgctccg caaggctgag 240  
catgcattct acttggaactg ggctgtccac tccttcagaa ttac 284

<210> 1121  
<211> 287  
<212> nucleic acid  
<213> Zea mays

<400> 1121

cgctcatgca caccgctgtg gatcttgtga acgagactaa gcttgacagc gagattaagt 60  
cttggettgc ttttgctgcc cagaagggtg ttgaggttga tgctcttgct aaggcattgg 120  
ctggtcacaaa ggatgaggct tacttcgcag caaatgctgc tgctcaggcc tcgaggaaat 180  
cctcaccocg tgtgaccaat gaagaagtcc agaaggctgc tgctgctctc aagggctctg 240  
accaccgccg tgggaccaac gttagtgtga gattggatgc ccagcag 287

<210> 1122  
<211> 290  
<212> nucleic acid  
<213> Zea mays

<400> 1122

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tggttgctt ttgctgccca gaaggttggt gaggttgatg ctcttgctaa ggcattggct 120  
ggcctaaaagg atgaggctta ctctgcagca aatgctgcag ctgaggctc gaggaaatcc 180  
tcaccccggtg tgaccaatga agaagtccag aaggctgctg ctgctctcaa gggctctgac 240

caccgccgtg cgaccaacgt tagtgctaga ttggatgcc agcagaagaa 290

<210> 1123  
<211> 344  
<212> nucleic acid  
<213> Zea mays

<400> 1123

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tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 120  
gctctccggt ttgcggttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 180  
caagcctccg atcancnacg gtgacgtgag ccgccccaac cccatgaccg ttttctggtc 240  
gaagacggca cagagcatga cgtctcgccc atgangggat gctgctggcc catcacaanc 300  
ctcaatngtc cttcgtccgg atgaccagcg agttgagntt ctac 344

<210> 1124  
<211> 261  
<212> nucleic acid  
<213> Zea mays

<400> 1124

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tgatgtgctt gtgcatggcg agcctgagag aaacgatatg gttgagtact tcggtgagca 120  
gctctccggt ttgcggttca ctgccaatgg ctgggtgcag tcttatgggt caaggtgcgt 180  
caagcctccg atcatctacg gtgacgtgag ccgccccaac cccatgaccg ttttctggtc 240  
gaagacggcg caaagcatga c 261

<210> 1125  
<211> 301  
<212> nucleic acid  
<213> Zea mays

<400> 1125

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aagctgctct ccgtgttccg anngggcgtc aaacncggcg cgggcatcgg ccccggtgtc 120  
tacgacatcc actccccag gatcccgtcg gctgaggaga tcgcccaccg catcgacaag 180

atgctggcgg tgctggacac caacatcctc tgggtgaacc ccgactgcgg cctcaagacc 240  
cgcaagtaca cggaggtcaa gcccgccctg accaacaatgg tctccgccgc taagtcctc 300  
c 301

<210> 1126  
<211> 264  
<212> nucleic acid  
<213> Zea mays

<400> 1126

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cccatatgtg ctactccaac ttcaacgnca tcatccactc gatcatcgac atggacgcgg 120  
acgtgntcac cattgagaac tcgcgggtccg acgagaagct gctctccgtg ttccgcgagg 180  
gcgtaagta cggcgccggc atcgccccg gtgtctacga catccnctcc ccagagatcc 240  
cgtcggctga ggagatcgcc gacc 264

<210> 1127  
<211> 277  
<212> nucleic acid  
<213> Zea mays

<400> 1127

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gacgagaagc tgctctccgt gttccgcgag ggcgtcaagt acggcgccgg catcgcccc 120  
ggtgtctacg acatccactc cccagagatc ccgtcggctg aggagatcgc cgaccgcctc 180  
gacaagatgc tggcgggtgct ggacaccaac atcctctggg tgaaccccga ctgcggcctc 240  
aagacccgca agtacacgga ggtcaagccc gcctgac 277

<210> 1128  
<211> 246  
<212> nucleic acid  
<213> Zea mays

<400> 1128

gggctgtcca ctctttcagg atcaccaact gcggagtcca ggacaccacc cagatccaca 60  
cccatatgtg ctactccaac ttcaacggac atcatccact ccatcatcga catggatgcc 120

gatgtgatca cgatcgagaa ctcccgatct gacgagaagc tactgtccgt cttccgtgag 180  
 ggtgtgaagt acggagctgg cattggccct ggtgtctacg acatccactc tcttaggatt 240  
 ccctcc 246

<210> 1129  
 <211> 270  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1129

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 agcctgagag aaacgctatg gttgagtact tcggtgagca gctctccggt tttgcgttca 120  
 ctgccaatgg ctgggtgcag tcttatgggt caaggtgctt caagcctccg atcatctacg 180  
 gtgacgtgag ccgccccaac cccatgaccg ttttctggtc gaagaggcgc agagcatgac 240  
 gtctcgccca atgaagggga tgctgactgg 270

<210> 1130  
 <211> 492  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1130

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 gcgggtccgac gagaagctgc tctccgtggt ccgcgagggc gtcaagtacg gcgcggggcat 120  
 cgcccccggt gtctacgaca tccactcccc caggatcccc tcggctgagg agatcgccga 180  
 ccgcatcgac aagatgctgg cgggtgctgga caccaacatc ctctgggtga accccgactg 240  
 cggcctcaag acccgcaagt acacggaggt caagcccgcc ctgaccaaca tgggtctccgc 300  
 cgctaaagct catccgcacc cagctcgcca gcgccaagtg agcacctttt ttttgccttt 360  
 ttcgtttccg aaganggcgt cntcgatgcc aatttgtttc caataaacan ggctgtnncc 420  
 cgccgttccg gttgtnancc gttcgggtgg aaggttanna attttcctga atttcccccc 480  
 aanggcggtt ng 492

<210> 1131  
 <211> 250  
 <212> nucleic acid



<213> Zea mays

<400> 1131

aactcaggag ggttcgccgt gaatacaagg cnaagaagat caccgaggac gaatacatca 60  
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tgcttgtgca tggagagcca gagagaaatg acatgggttg gtacttcggt gncnaattat 180  
ctggttttgc gttcaactgcc aacggatggg tgcaatccta tggatcacgc tgtgtgaagc 240  
cacccattat 250

<210> 1132

<211> 262

<212> nucleic acid

<213> Zea mays

<400> 1132

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gctgggtcaaa aggatgagge ttacttcgca gcaaagtctg ctgctcaggc ctcgaggaaa 120  
tcctcacccc gtgtgaccaa tgaagaagtc cagaaggctg ctgctgctct caagggctct 180  
gaccaccgcc gtgggaccaa cgttagtgtc agattggatg cccagcagaa gaagcttaac 240  
ctcccatcc ttcccaccac ta 262

<210> 1133

<211> 269

<212> nucleic acid

<213> Zea mays

<400> 1133

gctgctcagg cctcgaggaa atcctcaccc cgtgtgacca atgaagaagt ccagaaggct 60  
gctgctgctc tcaagggctc tgaccaccgc cgtgcgacca acgttagtgc tagattggat 120  
gccagcaga agaagcttaa cctcccatc cttccacca ctacaatcg ttcgttcna 180  
caaaccgttg agctcaggag ggtccgccgt gagtacaagg caaaaaagat ctctgaggag 240  
gaatacgtca ctgccatcaa ggaggaaat 269

<210> 1134

<211> 269

<212> nucleic acid

<213> Zea mays

<400> 1134

cgccgctgcg ccgtctcctc cgctcccgtc gtcgggaaag atggcgctccc acattgttgg 60  
 ataccctcgc atggggcccca agagggagct caagtttgca cttgagtctt tctgggatgg 120  
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180  
 gcagatggct gatgctggga tcaagtacat cccagcaac acccttctgt actatgacca 240  
 ggtccttgac accactgcc a tgctcggcg 269

<210> 1135

<211> 299

<212> nucleic acid

<213> Zea mays

<400> 1135

gcgatcaaga aggaagtcga ggatcttgag gctggtggta ttcaggttat ccaaatcgac 60  
 gaggctgcac tgagagaagg gctgccgntc cgcaaggctg agcatgcatt ctacttggac 120  
 tgggctgtcc actccttcag aattaccaac tgcgagatcc aggacaccac ccagatccac 180  
 acccacatgt gotantccaa cttcaacgac atcatccact cgatcatcga catggacgng 240  
 gacgtgatna ccattgagaa ntcgggtccg angggnagct gctctcngtg ttcngggag 299

<210> 1136

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1136

ctcaccocgt gtgaccaatg aagaagtcca gaaggctgct gctgctctca agggctctga 60  
 ccaccgcgct gggaccaacg ttagtgctag attggatgcc cagcagaaga agcttaacct 120  
 ccccatcctt cccaccacta caatcggttc gttcccacaa accgttgagc tcaggagggt 180  
 ccgccgtgag tacaaggcaa aaaagatctc tgaggaggaa tacgtcactg ccatcaagga 240  
 ggaaatcaac aaggtcgtta agcttcaaga agagct 276

<210> 1137

<211> 257

<212> nucleic acid

<213> Zea mays

<400> 1137

ccaactgcga gatccaggac accacccaga tccacacca catgtgctac tccaacttca 60  
acgacatcat ccaactcgatc atcgacatgg acgcggacgt gatcaccatc gagaactcgc 120  
ggtcgcgacga gaagctgctc tccgtgttcc gtgagggtgt caagtacggc gcggggcatcg 180  
gccccggtgt ctacgacatc cactcccccga ggatcccgtc ggccgaggag atcgccgacc 240  
gcatcgacaa gatgctg 257

<210> 1138

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 1138

gcttacttgc cagcaaattgc tgctgctcag gcctcgagga aatcctcacc ccgtgtgacc 60  
aatgaagaag tccagaaggc tgctgctgct ctcaagggct ctgaccaccg ccgtgggacc 120  
aacgttagtg ctagattgga tgcccagcag aagaagctta acctcccat ccttcccacc 180  
actacaatgc gttcgttccc acaaaccgtt gagctcagga gggccgcccg tgagtacaag 240  
gcaaaaaaga tctctgagga ggaatacgtc actgc 275

<210> 1139

<211> 295

<212> nucleic acid

<213> Zea mays

<400> 1139

agatggcaca gagcatgacc cctcgtccca tgaagggaat gttgactggc ccggtcacao 60  
tctgaactg gtcattcgtc aggaacgacc agcctagggt tgagacatgc taccaaatag 120  
ctcttgcaat caaaaaggag gttgaggatc ttgaggctgc tggattcag gtgatccaga 180  
togatgaggc agctctaagg gaggggtctgc cactacgcaa gtcagagcat gcattctacc 240  
tggactgggc tgtccactct ttcaggatca ccaactgcgg agtccaggac accac 295

<210> 1140

<211> 316

<212> nucleic acid

<213> Zea mays

<400> 1140

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gagacttact ttgctgatgt tctgtctgag tcttacaaga ccctaacatc cctgagcagt 120  
gtgactgctt atgggttttga tcttgtccgt ggaacccaaa ctcttgggct tgtcacgagt 180  
gctgggtttcc ctgctggaaa gtacctcttt gctgggtgtg tggatggacg caacatctgg 240  
gctgatgata ttgctacatc tctcagcact ctccagcttc ttgaggctgt tgttggaag 300  
gacaagcttg tcgtat 316

<210> 1141

<211> 445

<212> nucleic acid

<213> Zea mays

<400> 1141

ccactcttgt cctcgacctt gattctgaca aattggctgc attctctgct gcatacgcag 60  
aacttgaatc tgtactttct ggattgaatg tgcttgttga gacttacttt gctgatgttc 120  
ctgctgagtc ctacaagacc ctaacatctc tgagcagtgt gactgcctat ggttttgata 180  
ttgtccgtgg aacccaaact cttgggcttg tcacgagtgc tggtttccct gctggaaagt 240  
acctctttgc tgggtgttgg gatggacgca acatctgggc tgatgatctt gctacatctc 300  
taacactctt cagtctcttg aggctgttgt tgggaangac aagcttgtcg atcaacttnc 360  
tgctcgctca tgcacaccgn tgtggatctt gngaaccaga ctnagcttga cagcagatna 420  
atctggcttg ctttgcctgc anaag 445

<210> 1142

<211> 301

<212> nucleic acid

<213> Zea mays

<400> 1142

gccaaagcgg ttgttgaggt tgatgctctt gctaaggcat tggctggtca aaaggatgag 60  
gcttacttgc cagcaaatgc tgcagctcag gctcagga aatcctcacc ccgtgtgacc 120  
aatgaagaag tccagaaggc tgctgctgct ctcaagggt ctgaccaccg ccgtgcgacc 180

aacgttagtg ctagattgga tgcccagcag aagaagctta acctcccat cttcccacca 240  
tacaatcggg tcgttccac aaaccgttga gtcaggagg gtccgccgtg agtacaaggc 300  
a 301

<210> 1143  
<211> 277  
<212> nucleic acid  
<213> Zea mays

<400> 1143

gtggtattca gggtatccaa atcgacgagg ctgcactgag agaagggctg ccgctccgca 60  
aggctgagca tgcattctac ttggactggg ctgtccaact cttcagaatc accaactgcg 120  
agatccagga caccaccng atccacaccc acatgtgcta ctccaacttc aacgacatca 180  
tcactcgat catcgacatg gacgaggacg tgatcaccat cgagaatcgc ggtccgacga 240  
gaagctgctc tccgtgttcc gtgaggtgtc aagtacg 277

<210> 1144  
<211> 281  
<212> nucleic acid  
<213> Zea mays

<400> 1144

cgccgtgcg ccgtctctc cgctcccgtc gtccggaaag atggcgtccc acattgttgg 60  
ataccctcgc atggggccca agagggagct caagtgtgca cttgagtctt tctgggatgg 120  
gaagagcacc gctgaggatc tggagaangt tgctaccgac ctgagggccca gtatctggaa 180  
gcagatggct gatgctggga tcaagtacat cccagcaac actttctcgt actatgacca 240  
ggtcctttga caccactgcc atgctcggcg ctgtccccga a 281

<210> 1145  
<211> 247  
<212> nucleic acid  
<213> Zea mays

<400> 1145

gccgtgctac cactgttgct gctagattgg atgctcagca gaaaaagctc aaccttcctg 60  
tccttccac aaccacaatt ggttcattcc ctgagactgt ggaactcagg agggtcgcgtc 120

gtgagtacaa ggcaaagaag atctccgagg aggaatacat cagtgccatc aaggaagaaa 180  
 tcagcaaggt tgtcaagatc caagaggagc ttgacattga tgtgcttgtg catggagagc 240  
 cagagag 247

<210> 1146  
 <211> 268  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1146

tgctgactgg cccagtcaca atcctcaact ggtccttcgt ccggaatgac cagccgaggt 60  
 ttgagacttg ctaccagatc gctcttgcca tcaagaagga agtcgaggat cttgaggctg 120  
 gtggtattca ggttatccaa atcgacgagg ctgcaactgag agaagggctg ccgctccgca 180  
 aggctgagca tgcattctac ttggactggg ctgtccactc cttcagaatt accaactgcg 240  
 agatccagga caccacccag atccacac 268

<210> 1147  
 <211> 290  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1147

cggnacggtg gcnacctatc atcgccgctg cgcggtctcc tccgctcccg tcgtccggaa 60  
 agatggcgtc ccacattggt ggataccctc gnntggggccc caagagggag ctcaagtttg 120  
 cacttgagtc tttctgggat ggggaagagca ccgctgagga tctggagaag gttgctaccg 180  
 acctcagggc cagtatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 240  
 acactttctc gtactatgac caggtccttg acaccactgc catgctcggc 290

<210> 1148  
 <211> 269  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1148

caaggagatc atgccccagg gtgacaaacg aggaggtcca gaaggctgca gctgctttga 60  
 ggggatctga ccaccgccgt tctaccactg tttctgctag attggatgct cagcagaaaa 120

agctcaacct tctgtcctt cccacaacca caattgggtc attccctcag actgtggaac 180  
 tcaggagggt tcgccgtgaa tacaaggcaa agaagatcac cgaggacgaa tacatcagtg 240  
 ccatcaagga agaaatcagc aaggttgtc 269

<210> 1149  
 <211> 280  
 <212> nucleic acid  
 <213> Zea mays

<400> 1149

caagaagagc ttgacatnna tgncttgtn catngcgagc ctgagagaaa cgatatgggtt 60  
 gagnacntcg gtgagcagct cncgggtttt gcgttcacng ccaatggctg ggtgcagtct 120  
 tatgggtcaa ggtgogtcaa gcctccgac atctacgggtg acgtgagcng ccccaacccc 180  
 atgaccgttt tcnngtgcgaa gacngcgag agcatgacgt ctgcccgaat gaaggggatg 240  
 ctganggccc agtcacaatc ctcaacnggt ccttcgtccg 280

<210> 1150  
 <211> 323  
 <212> nucleic acid  
 <213> Zea mays

<400> 1150

gagaactcgc ggtccgacga gaagctgctc tccgtgttcc gtgaggggtg caagtacggc 60  
 gcgggcatcg gccccggtgt ctacgacatc cactcccca ggatcccgtc ggccgaggag 120  
 atgcgcgacc gcacgcgaaa gatgctggcc gtgctggaca ccaacatcct ctgggtgaac 180  
 cccgactgcg gcctcaagac ccgcaagtac acggaggtca agcccgccct gaccaacatg 240  
 gtctccgctg ctaagtcatt cgcacccagc tcgccagcgc caagtgcgca ctttttttgc 300  
 ctttttctgt tccgaggagg ggt 323

<210> 1151  
 <211> 325  
 <212> nucleic acid  
 <213> Zea mays

<400> 1151

caccacccac atctcagttc accccggcgt cgtcctcttc ccccggcgt actctccgc 60

tccacgttcc aaaggaaaga tggcgtccca tattgttga taccctcgca tgggccccaa 120  
gagagagctc aagtttgcct tggagtcttt ctgggatggg aagagcagcg ccgatgattt 180  
agagaaagtt gcctctgacc tcaggtctag catctggaag caaatgtcag aagctgggat 240  
caagtacatt cccagcaaca ccttctcgta ctatgaccag gtccttgata ccacggccat 300  
gotttggtgtg tcccagagcg ctact 325

<210> 1152  
<211> 272  
<212> nucleic acid  
<213> Zea mays

<400> 1152

ggacgtggac ttgcgatggg cccaagagg gagctcaagt ttgcacttga gtcantctgg 60  
gatgggaaga gcaccgctga ggatctggag aaggttgcta ccgacctcag ggccagtatc 120  
tggangcaga tggctgatgc tgggatcaag tacatcccca gcaacacctt ctctactat 180  
gaccaggtcc ttgacaccac tgccatgctc ggcgctgtcc ccgaacgcta ctcatggact 240  
ggaggagaga ttgggtttga cacctacttc tc 272

<210> 1153  
<211> 232  
<212> nucleic acid  
<213> Zea mays

<400> 1153

gggctgtcca ctcttcaga atcaccaact gcgagatcca ggacaccacc cagatccaca 60  
cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgac atggacgcgg 120  
acgtgatcac catcgagaac tcgcggtccg acgagaagct gctctccgtg ttccgtgagg 180  
gtgtcaagta cggcgcgggc atcggtcccg gtgtctacga catccactcc cc 232

<210> 1154  
<211> 285  
<212> nucleic acid  
<213> Zea mays

<400> 1154

atcgacgagg ctgcactgag agaagggctg ccgctccgca aggctgagca tgcattctac 60



ttggactggg ctgtccactc cttcagaatc accaactgcg aaatccagga caccacccag 120  
atccacaccc acatgtgcta ctccacttca acgacatcat ccaactcgatt nccgacatgg 180  
acgcggacgt gatcaccatc gagaactcgc ggtccgacga gaagctgctc tccgtgttcc 240  
gtgaggtgtc aagtacggcg cgggcacgcg ccccggtgtc tacga 285

<210> 1155  
<211> 327  
<212> nucleic acid  
<213> Zea mays

<400> 1155

cccggtctga cgagaagcta ctgtccgtct tccgtgaggg tgtgaagtac ggagctggca 60  
ttggccctgg tgtctacgac atccacttct cctaggattc cctccaccga ggagatcgca 120  
gaccgcgtcg agaagatgct cgcggtgctc gacaccaaca tcctctgggt gaaccctgac 180  
tgtggtctca agacacgcaa gtacacggag gtcaagcccg ccctgaccaa catggtctcg 240  
gccaccaagc tcattccgac ccagcttgcc agcgcgaaat gagtcgtttg atagtccatg 300  
gtctgatagc gcggaatgag ccagttt 327

<210> 1156  
<211> 284  
<212> nucleic acid  
<213> Zea mays

<400> 1156

cgatgacacc attgaganct cgcggtccga cgagaagctg ctctccgtgt tccgcgaggn 60  
gcgtcaagta cggcgcgggc atcgccccg gtgtctacna catccactcc ccaggatcc 120  
cgtcggctga ggagatcgcc gaccgcatcg acaagatgct ggcggtgctg gacaccaaca 180  
tcctctgggt gaaccccgac tgcggcctca agaccgcaa gtacacggag gtcaagcccg 240  
ccctgaccaa catggtctcc gccgctaagc tcattccgac ccag 284

<210> 1157  
<211> 293  
<212> nucleic acid  
<213> Zea mays

<400> 1157

gtcttgggct tgcttttgct gcccagaagg ttgttgaggt tgatgctctt gctaaggcat 60  
 tgggctggtc aaaaggatga ggcttacttc gcagcaaagt ctgctgctca ggctcgagg 120  
 aaatcctcac ccggtgtgac caatgaagaa gtccagaagg ctgctgctgc tctcaagggc 180  
 totgaccacc gccgtgggac caacgttagt gctagattgg atgccagca gaagaagctt 240  
 aacctcccca tcttccac cactacaatc ggttcgttcc cacaaccgt tga 293

<210> 1158  
 <211> 309  
 <212> nucleic acid  
 <213> Zea mays

<400> 1158

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 ataccctcgc atgggccccca agagggagct caagtttgca cttgagtctt tctgggatgg 120  
 gaagggcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180  
 gcagatggct gatgctggga tcaagtacat cccagaanc actttctcgt actatgacca 240  
 agtcttgaca ccatgccatg ctggcgctg tccccgaac ttactcatgg actggaggag 300  
 agatggatt 309

<210> 1159  
 <211> 462  
 <212> nucleic acid  
 <213> Zea mays

<400> 1159

cgtgcggtgn tccgcgaggg cgtcaagtac ggcgcnggca tcggccccgg tgtctacgac 60  
 ntccactccc ccaggatccc gtcggctgag gagatcgccg accgcatcga caagatgctg 120  
 gcggtgctgg acaccaacat cctctgggtg aaccccgact gcggcctcaa gaccgcaag 180  
 tacacggagg tcaagccgc cctgaccaac atgggtctccg ccgctaagct catccgcacc 240  
 cagctcgcca gcgccaagtg agcacctttt ttttgctttt ttcgtttccg aggagggccg 300  
 tcgtcgatgc caatttgttt ccaataaaca gggcttgctc ccgccgttct gttgtactcc 360  
 gtctgtggtt aggttaagta gttttcttga tctcgcccc acgcggtacc tgnittacta 420  
 ctctctggtt ggtgggttct gaagcaaagt tnnccgttta ct 462

<210> 1160  
 <211> 231  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1160  
  
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 ggctgtccac tccttcagaa tcaccaactg cgagatccag gacaccaccc agatccacac 120  
 ccacatgtgc tactccaact tcaacgacat catccactcg atcatcgaca tggacgcgga 180  
 cgtgatcacc atcgagaact cgcgggtccga cgagaagctg ctctccgtgt t 231

<210> 1161  
 <211> 275  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1161  
  
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 tgagacatgc taccaaatag ctcttgcaat caaaaaggag gttgaggatc ttgaggcnnc 120  
 nggtattcag gtgatccaga tcgatgaggc agctctaagg gagggctctgc cactacgcaa 180  
 gtcagagcat gcattctacc tggactgggc tgtccactct ttcaggatca ccaactgcgg 240  
 agtccaggac accaccaga tccacaccca catgt 275

<210> 1162  
 <211> 290  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1162  
  
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 tggagatgac caagtggttt gacaccaact accactttan tgtccctgga nttgggcca 120  
 aacaccaagt tctcctatgc ttctcacaag gctgttaacg aatataagga ggctaaggcg 180  
 ctcggtgttg ataccgtgcc agtacttggt ggaccagtct cgtacctgtt gctctcaaag 240  
 cctgccaaagg gtgtggagaa gggattcnct cttctttccc ttcttagcag 290

<210> 1163

<211> 342  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1163  
  
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 gacgtctcng ccaatgaaag gngatgctga ctggcccagt ncacaatcct caactgggtcc 120  
 ttcgtccgga atgaccagcc gaggtttgag acttgctacc agatcgtctt tgcgatcaag 180  
 aaggaagtcg aggatctcga ggctgggtgg attcagggtta tccaaatcga cgaggctgca 240  
 ctgagagaag ggctgccgt cgcgaaggct gagcatgcat ctacttgga tgggctgtcc 300  
 atccttcaga atcaccaatg cgnagaccag gacaccaccc ag 342

<210> 1164  
 <211> 264  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1164  
  
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 ggatgctgac tggcccagtc acaatcctca actggctcct cgtccggaat gaccagccga 120  
 ggtttgagac ttgctaccag atcgtctctt cgatcaagaa ggaagtcgag gatcttgagg 180  
 ctggtggtat tcaggttatc caaatcgacg aggctgcact gagagaaggg ctgccgctcc 240  
 gcaaggctga gcatgcattc tact 264

<210> 1165  
 <211> 298  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1165  
  
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 gctagattgg atgctcagca gacaaagctc aaccttnctg tccttccac aaccacaatt 120  
 gnttcattcc ctcagactgt ggaactcagg agggttcgcc gtgaatacaa ggcaaagaag 180  
 atcaccgagg acgaaaanat cagtgccatc aaggaagaaa ttcagcaagg ttgtcaagat 240  
 ccaagaggag ntgacattga tgtgcttgtg catggagagc cagagagaat gacatggt 298

<210> 1166  
 <211> 226  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1166  
  
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 cgggtgagcaa ttatctgggtt ttgcgttcac tgccaacgga tgggtgcaat cctatggatc 120  
 acgctgtgtg aagccaccca ttatctacgg tgatgtcagc cggccgaacc ccatgactgt 180  
 tttctgggtcc aagatggcac agagcatgac cctcgtccc atgaag 226

<210> 1167  
 <211> 254  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1167  
  
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 gaagtgcagg atcttgaggc tgggtgtatt caggttatcc aaatcgacga ggctgcactg 120  
 agagaagggc tgccgctccg caaggctgag catgcattct acttggactg ggctgtccac 180  
 tccttcagaa ttaccaactg cgagatccag gacaccaccc agatccacac ccacatgtgc 240  
 tactccaact tcaa 254

<210> 1168  
 <211> 251  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1168  
  
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 gcctccgac atctacggtg acgtgagccg cccaacccc atgaccgttt tctggtcgaa 120  
 gaggcgcaga gcatgacgtc tcgccaatg aaggggatgc tgactggccc agtcacaatc 180  
 ctcaactggt ccttcgtccg gaatgaccag ccgaggtttg agacttgcta ccagatcgct 240  
 cttgcgatca a 251

<210> 1169

<211> 279  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1169  
  
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 ggataccctc gcatggggccc caagagggag ctcaagtttg cacttgagtc tttctgggat 120  
 gggaagagca ccgctgagga tctggagaag gttgcnacng aacttnaggg ccagtatctg 180  
 gaagcagang gctgatgctg ggatcaagta catccccagc aacactttct cgtactatga 240  
 ccaggtcctt gacacnactg ccatgctggg gctgggtccc 279

<210> 1170  
 <211> 268  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1170  
  
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 tgaccgtttt ctggctgaag aggcgcagan catgacgtct cgcccaatga aggggatgct 120  
 gactggccca gtcacaatcc tcaactggtc cttcgtccgg aatgaccagc cgaggtttga 180  
 gaattgctac cagatcgctc ttgcgatcaa gaaggaagtc gaggatctcg aggctgggtgg 240  
 tattcagggt atccaaatcg acgaggct 268

<210> 1171  
 <211> 90  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1171  
  
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 tcntctgggt gaaccccgac tgcngcctca 90

<210> 1172  
 <211> 254  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1172

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 gcccgaagag ggagctcaag ttgacattg agtctttctg ggatgggaag agcaccgctg 120  
 aggatctgga gaaggttget accgacctca gggccagtat ctggaagcag atggctgatg 180  
 ctgggatcaa gtacatcccc agcaacacct tntcgtaata tgaccaggtc cttgacacca 240  
 tgccatgctc ggnt 254

<210> 1173  
 <211> 251  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1173

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 ccaactgtttc tgctagattg gatgctcagc agaaaaagct caaccttcct gtccttccca 120  
 caaccacaat tggttcattc cctcagactg tggaactcag gagggttcgc cgtgaatata 180  
 aggcaaagaa gatcaccgag gacgaatata tcagtgnat caaggaagaa atcagcaagg 240  
 ttgtcaagat c 251

<210> 1174  
 <211> 246  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1174

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 aggggtgtgaa gtacggagct ggcattggcc ctgggtgtcta cgacatccac tctcctagga 120  
 ttccctccac cgaggagatc gcagaccgcg tcgagaagat gctcgccgtg ctcgacacca 180  
 acatcctctg ggtgaacct gactgtggtc tcaagacacg caagtacacg gaggtcaagc 240  
 ccgccc 246

<210> 1175  
 <211> 258  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1175

gaacgatcag cctaggtttg agacatgcta ccaaatagct cttgcaataa aaaaggaggt 60  
 tgaggatctt gaggtgctg gtattcaggt gatccagatt gatgaggcag ctctaaggga 120  
 ggggtctgcca ctgcgcaagt cagagcatgc attctacctg gactgggctg tccactcttt 180  
 cagaatcacc aactgcggag tccaggacac caccagatc cacaccata tgtgctactc 240  
 caacttcaac gacatcat 258

<210> 1176  
 <211> 276  
 <212> nucleic acid  
 <213> Zea mays

<400> 1176

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 tgctgctctc aagggtctctn aanaccgctg nngnaccaa gttagtgtga gattggatgc 120  
 ccagcagaag aagcttaacc tccccatcct tcccaccact acaatcggtt cgttcccaca 180  
 aaccgttgag ctgaggaggg tccgccgtga gtacaaggca aaaaagatct ctgaggagga 240  
 atacgtcact gccatcaagg aggaaatcaa caaggt 276

<210> 1177  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays

<400> 1177

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 atctgggctg atgatcttgc tacatctctc agcactctcc agtctcttga ggctgttggt 120  
 gggaaggaca agcttgctgt atcaacttcc tgctcgctca tgcacaccgc tgtggatctt 180  
 gtgaacgaga ctaagcttga cagcgagatt aagtcttggc ttgctttttg cccagaaggt 240  
 tgttgagggt gatgctcttg ctaaggcatt ggctgggcca aaagatgagc ttacttc 297

<210> 1178  
 <211> 283  
 <212> nucleic acid  
 <213> Zea mays

<400> 1178



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gttgcctctca aagcctgccca aggggtgtgga gaagggattc cctcttcttt cccttcttag 120  
cagcatcctc ccagtctaca aggaggtcat tgctgagttg aaggcagctg gggcttcatg 180  
gattcagttt gatgagccca ctcttgtcct cgaccttgat tctgacaaat tggctgcatt 240  
ctctgctgca tacgcagaac ttgaatctgt actttctgga ttg 283

<210> 1179  
<211> 84  
<212> nucleic acid  
<213> Zea mays

<400> 1179

ancttctngt actatgacca ggtccttgan accagtncca tgctcggcgc tgtccccgaa 60  
cgctaccatg gactggacga gaga 84

<210> 1180  
<211> 315  
<212> nucleic acid  
<213> Zea mays

<400> 1180

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tgcatacgca gaacttgaat ctgtactttc tggattgaat gtgcttggtg agacttactt 120  
tgctgatgtt cctgctgagt cctacaagnc cctaacatcc ctgagcagtg tgactgctta 180  
tggttttgat cttgtccgtg gaacccaaac tcttgggctt gtcacgagtg ctggtttccc 240  
tgctggaaaag tacctctttg ctggtgttgt ggatggacgc aacatctggg ctgatgatct 300  
tgctacatct ctgag 315

<210> 1181  
<211> 267  
<212> nucleic acid  
<213> Zea mays

<400> 1181

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caatgaaggg gatgctgact ggcccagtca caatcctcaa ctggctcctc gtccggaatg 120

accagccgag gtttgagact tgctaccaga tcgctcttgc gatcaagaag gaagtcgagg 180  
 atctcgaggc tgggtggtatt caggttatcc aaatcgacga ggctgcactg agagaagggc 240  
 tgccgtccgc aaggctgagc atgcatt 267

<210> 1182  
 <211> 249  
 <212> nucleic acid  
 <213> Zea mays

<400> 1182

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 ctccacagag gagatcgag accgcgtcga gaagatgctc gccgtgctcg acaccaacat 120  
 cctctgggtg aaccctgact gtgggtctcaa gacacgcaag tacacggagg tcaagcccgc 180  
 cctgaccaac atgggtctcg ccaccaagct catccgcacc cagcttgcca gcgcgaaatg 240  
 aggtcgttt 249

<210> 1183  
 <211> 259  
 <212> nucleic acid  
 <213> Zea mays

<400> 1183

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 ccacattggt ggataccctc gcatgggccc caagagggng ctcaagtttg cacttgagtc 120  
 tttctgggat gggaagagca ccgctgagga tctggagaag gttgctaccg acctcagggc 180  
 cagtatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca acactttctc 240  
 gtactatgac caggtcctt 259

<210> 1184  
 <211> 296  
 <212> nucleic acid  
 <213> Zea mays

<400> 1184

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tgcattctac ctggactggg ctgtccactc tttcaggatc accaactgcg gagtccagga 180  
caccacccag atccacaccc acatgtgcta ctccaacttc aacgacatca tccactccat 240  
ccatcgacat gggatgccga tgtgatcacg atcgagaact cccggtctga ngagaa 296

<210> 1185  
<211> 218  
<212> nucleic acid  
<213> Zea mays  
<400> 1185

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cccacatgtg ctactccaac ttcaacgaca tcatccactc gatcatcgac atggacgcgg 180  
acgtgatcac catcgagaac tcgcggtccg acgagaag 218

<210> 1186  
<211> 261  
<212> nucleic acid  
<213> Zea mays  
<400> 1186

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gtccacatt gttggatacc ctgcgatggg cccaagagg gagtcaagt ttgcacttga 120  
gtttttctgg gatgggaaga gcaccgctga ggatctggag aaggttgcta ccgacctcag 180  
ggccagtatc tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaacacttt 240  
ctcgtactat gaccaggtcc t 261

<210> 1187  
<211> 282  
<212> nucleic acid  
<213> Zea mays  
<400> 1187

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gagtctttct gggatgggaa gagcaccgct gaggatctgg agaaggttgc taccgacctc 120  
agggccagta tctggaagca gatggctgat gctgggatca agtacatccc cagcaacact 180

ttctcgtact atgaccaggt ccttgacaac atgccatgct cggcgtgtcc ccgaacgtta 240  
ctcatggatn gaggagagat ggattgacaa ctattctcca tg 282

<210> 1188  
<211> 298  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1188

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ctggtcgaag aggcacagag catgacgtct cgcccaatga aggggatgct gactggccca 120  
gtcacaatcc tcaatggtcc ttctgtccga atgaccagcc gaggtttgag acttgctacc 180  
agatcgctct tgcgatcaag aaggaagtcg aggatcttga ggctgggtgg attcaggtta 240  
tccaaatoga cgaggctgcc actgagagaa gggctgccgt ccgcaaggct gagcatca 298

<210> 1189  
<211> 228  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1189

cccagatcca caccacatg tgctactcca acttcaacga catcatccac tncgatcatc 60  
gacatggacg cggacgtgat caccatcgag aactcgcggc ccgacgagaa gctgctctcc 120  
gtgttccgtg aggggtgtcaa gtacggcgcg ggcacggcc ccggtgtcta cgacatccac 180  
tccccagga tcccgtcggc cgaggagatc gccgaccga tcgacaag 228

<210> 1190  
<211> 280  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1190

cacaattggt tcattccctc agactgtgga actcaggagg gttccgccgt gaatacaagg 60  
caaagaagat caccgaggac gaatacatca gtgccatcaa ggaagaaatc agcaaggncg 120  
tcaagatcca agaggagctt gacattgatg tgcttgtgca tggagagcca gagagaaatg 180  
acatggttga gtacttcggt gagcaattat ctggtttgcg ttcactgcca acggatgggt 240

gcaatcctat ggatcacgct tgtgtgaagc caccattat 280

<210> 1191  
<211> 290  
<212> nucleic acid  
<213> Zea mays

<400> 1191

ttgggaagga caagcttgct gtatcaactt cctgctcgt catgcacacc gctgtggatc 60  
ttgtgaacga gactaagctt gacagcgaga ttaagtctng gcttgctttt gctgcccaga 120  
aggttggtga gggtgatgct cttgctaagg cattggctgg tcaaaaggat gaggettact 180  
tcgcagcaaa tgctgctgct caggcctcga ggaaatcctc acccgtgtg accatgaaga 240  
agtccagaag gctgctgctg ctctcaaggg tctgaccacc gccgtgggac 290

<210> 1192  
<211> 243  
<212> nucleic acid  
<213> Zea mays

<400> 1192

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gatcttgagg ctgggtggtat tcagggttatc caaatcgacg aggctgcact gagagaaggg 120  
ctgccgctcc gcaagggtga gcatgcattc tacttggtact gggctgtcca ctccttcaga 180  
attaccaact gcgagatcca ggacaccacc cagatccaca cccacatgtg ctactccaat 240  
tca 243

<210> 1193  
<211> 223  
<212> nucleic acid  
<213> Zea mays

<400> 1193

atggacgcag atgtgatcac catcgagaac tcccgggtccg acgagaagct actgtccgtg 60  
ttccgcgagg gtgtgaagta cggagctgga atcgggtccag gtgtctacga catccactcg 120  
cccaggatcc cctccaccga ggagatcgcg gaccgctca acaagatgct cgccgtgctc 180  
gacaccaaca tcctctgggt gaaccctgac tgcggtctca aga 223

<210> 1194  
 <211> 246  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1194  
  
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 tgagacttgc taccagatcg ctcttgcat caagaaggaa gtcgaggatc tcgaggctgg 120  
 tggatattcag gttatccaaa tcgacgaggc tgactgaga gaagggtgc cgctccgcaa 180  
 ggctgagcat gcattctact tggactgggc tgtccactcc ttnagantca ccacngggag 240  
 atccng 246

<210> 1195  
 <211> 232  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1195  
  
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 caaggctgan catgcattct acttgactn ggctgtccac tccttcagaa ttaccaactg 120  
 cgagatccag gacaccaccc agatccacac ccacatgtgc tactccaact tcaacgacat 180  
 catccactcg atcatcgaca tggacgcgga cgtgatcacc attgagaact cg 232

<210> 1196  
 <211> 248  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1196  
  
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 gagcctgaga gaaacgatat ggttgagtac ttcggtgagc agctctccgg ttttgcggtc 120  
 actgccaatg gctgggtgca gtcttatggg tcaagggtgcg tcaagcctcc gatcatctac 180  
 ggtgacgtga gccgccccaa ccccatgacc gttttctggt cgaagaggcg cagagcatga 240  
 cgtctcgc 248

<210> 1197

<211> 300  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1197  
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 cttgaggctg ttgttgggaa ggacaagctt gtcgtatcaa cttcctgctc gctcatgcac 120  
 accgctgtgg atcttgtgaa cgagactaag cttgacagcg agattaagtc ttggcttgct 180  
 tttgctgccc agaaggttgt tgaggttgat gctcttgcta aggcattggc tggcmetaag 240  
 gatgaggcta cttcgcagca aatgctgctg ctcaggcctc gaggaatcc tcaccccgctg 300

<210> 1198  
 <211> 233  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1198  
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 tgcattctac ctggactggg ctgtccactc tttcaggatc accaactgcg gagtccagga 120  
 caccacccag atccacaccc acatgtgcta ctccaacttc aacgacatca tccactccat 180  
 catcgacatg gatgccgatg tgatcacgat cgagaatccc ggtctgacga gaa 233

<210> 1199  
 <211> 256  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1199  
 gggaacgacc agccgaggtt tgagacttgc taccagatcg ctcttgcgat caagaaggaa 60  
 gtcgaggatt cttggaggct ggtggtattc aggttatcca aatcgacgag gctgcactga 120  
 gagaagggct gccgctccgc aaggctgagc atgcattcta cttggactgg gctgtccact 180  
 ccttcagaat taccaactgc gagatccagg acaccaccca gatccacacc cacatgtgct 240  
 actccaactt caacga 256

<210> 1200  
 <211> 297  
 <212> nucleic acid

<213> Zea mays

<400> 1200

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aggattccct ccaccgagga gatcgagac cgctcgaga agatgctgc cgtgctcgac 120  
accaacatcc tctgggtgaa ccctgactgt ggtctcaaga cacgcaagta cacggaggtc 180  
aagccccccc tgaccaacat ggtctcggcc accaagctca tccgcaccca gcttgccagc 240  
gcgaaatgag tcgtttgata gctccatggt ctgatagcgc ggaatgagcc agtttgt 297

<210> 1201

<211> 260

<212> nucleic acid

<213> Zea mays

<400> 1201

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acattgttgg ataccctcgc atgggccccca agaggagct caagtttgca cttgagtctt 120  
tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggccca 180  
gtatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac ancttctcgt 240  
actatgacca ggtcttgaca 260

<210> 1202

<211> 261

<212> nucleic acid

<213> Zea mays

<400> 1202

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ccccaggatc ccgtcggctg aggagatcgc cgaccgcatc gacaagatgc tggcgggtgct 120  
ggacaccaac atcctctggg tgaaccccga ctgcggcctc aagacncgca agacccacgg 180  
aggtcaagcc cncctgacn aacatggtct ccgccgctaa gctcatncgc acccagntcg 240  
ccagcgccaa gtgagcactt t 261

<210> 1203

<211> 246

<212> nucleic acid



<213> Zea mays

<400> 1203

cccttgccaa ggctttggca ggccaaaagg atgaggtcta ctttgcagcc aatgctgctg 60  
ctnnggcctc aaggagatca tcgcccaggg tgacaaaacga ggaggtccag aaggctgcag 120  
ctgctttgag gggatctgac caccgccgtt ctaccactgt ttctgctaga ttggatgctc 180  
agcagaaaaa gctcaacctt cctgtccttc ccacaaccac aattggttca ttccctcaga 240  
ctgtgg 246

<210> 1204

<211> 266

<212> nucleic acid

<213> Zea mays

<400> 1204

gggacntggt ccagtgggta ncgcggacgt gatcaccatt ganaactcgc ggtccgaagn 60  
gaagctgctc tccgtgttcc gcnagggcgt caagtacggc gcgggcatcg gccccggtgt 120  
ctacgacatc cactccccca ggatcccgtc ggctgaggat atcgccgacc gcatcgacaa 180  
gatgctggcg gtgetggaca ccaacatcct ctgggtgaac cccgactgcg gcctcaagac 240  
ccncaagtac acggaggctc agcccc 266

<210> 1205

<211> 302

<212> nucleic acid

<213> Zea mays

<400> 1205

ccccaacccc atgaccgttt tcnggtcgaa gacggcacag agcatgacgt ctgcccgaat 60  
gaaggggatg ctgactggcc cagtcacaat cctcaactgg tccttcgtcc ggaatgacca 120  
gcogangttt gagacttgct accagatcgc tcttgcgatc aagaaggaag tcgaggatct 180  
tgagtntggt ggtattcagg ttatccaaat cgacgaggct gcactgagan aagggcnnnn 240  
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ag 302

<210> 1206

<211> 319  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1206  
  
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 caaccccatg accgttttct ggtcgaagac ggcgagagc atgacgtctc gcccaatgaa 120  
 ggggatgctg actggcccag tcacaatcct caactgggtc ttcgtccgga atgaccagcc 180  
 gaggtttgag acttgctacc agatcgctct tgcgatcaag aaggaatcga ggactcgagg 240  
 ctggtggtat cagttaccaa ncgagagtgc ntgagagaag gtgcgtcgca gntgacnnat 300  
 ctattgatgg tgtcatctc 319

<210> 1207  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1207  
  
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 tgggtgttg gatggaagca acatctgggc tgatgatctt gctacatctc tcagcantct 120  
 ccagtctctt gaggtgttg ttgggaagga caagcttgct gtatcaactt cctgctcgct 180  
 catgcacacc gctgtggatc ttgtgaacga gactaagctt gacagcgaga ttaagtcttg 240  
 gcttgctttt gctgcccaga angttgttga gttgatgtct tgctaaaggc attgctggt 299

<210> 1208  
 <211> 315  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1208  
  
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 tggtggtca aaaggatgag gcttacttcg cagcaaagtc tgcngtcagg cctcgaggaa 180  
 atcctcacc cgtgtgacca atgaagaagt ccagaaggct gntgctgctc tcaagggctc 240  
 tgaccaccgc cgtgggacca acgttagtgc tagatggatg cccagcngtn agnagcttaa 300

cctccccanc ctccc

315

<210> 1209  
<211> 421  
<212> nucleic acid  
<213> Zea mays

<400> 1209

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accgntgtgg atcttgtgaa cgagactaat cttgacagcg agattaagtc ttggcttgct 120  
tttgctgccc agaaggttgt tgaggttgat gctcttgcta aggcatggc tggcctaaaag 180  
gatgaggctt acttcgcagc aaatgctgct gctcaggcct cgaggaaatc tcaccccgtn 240  
tgaccaatga anaantccan aaggctgctg ctgctctcaa gggctctgac caccggcgtn 300  
ggaccaacgt tagtgctaga ttggatgccc acaaangaaa cttaacctcc cattcttcca 360  
ccacttcaat tggttcgtnn ccacaaaccc gttgagctca agaagggtccg ccgtgagtac 420  
a 421

<210> 1210  
<211> 261  
<212> nucleic acid  
<213> Zea mays

<400> 1210

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cgaccgcacg gacaagatgc tggccgtgct ggacaccaac atcctctggg tgaaccccca 120  
ctgcggcctc aagaccgcga agtacacgga ggtcaagccc gccctgacca acatggtctc 180  
cgctgctaag ctcatcgca cccagctcgc cagcgccaag tgagcacctt tttttgcctt 240  
tttcgtttcc gaggagggcg t 261

<210> 1211  
<211> 239  
<212> nucleic acid  
<213> Zea mays

<400> 1211

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gtgaacgaga ctaagcttga cagcgagatt aagtcttggc ttgcttttgc tgcccagaag 120  
 gttgttgagg ttgatgctct tgctaaggca ttggctggtc aaaaggatga ggcttacttc 180  
 gcagcaaatg ctgctgctca ggctcgagg aaatcctcac cccgtgtgac caatgaaga 239

<210> 1212  
 <211> 236  
 <212> nucleic acid  
 <213> Zea mays

<400> 1212

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 gggaagagca ccgctgagga tctggagaag gttgctaccg acctcagggc cagtatctgg 180  
 aagcagatgg ctgatgctgg gatcaagtac atcccagca acactttctc gtacta 236

<210> 1213  
 <211> 276  
 <212> nucleic acid  
 <213> Zea mays

<400> 1213

cccacatctc agttcacccc ggcgctgctc tcttcccccg gcgctactct cccgctccac 60  
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 agctcaagtt tgccttggag tctttctggg atgggaagag cagcgccgat gatttagaga 180  
 aagttgcctc tgacctcagg tctagcatct ggaagcaaat gtcagaagct gggatcaagt 240  
 acattcccag caacaccttc tcgtactatg accagg 276

<210> 1214  
 <211> 231  
 <212> nucleic acid  
 <213> Zea mays

<400> 1214

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 tctggctcaa gacggcgcag agcatgacgt ctgcaccaat gaaggggatg ctgantggcc 120  
 cagtcacaat cctcaactgg tcttctgctc ggaatgacca gccgaggttt gagan ttgcn 180

ancagatcgc tcttgcgac aagaaggaag tngaggatct cgangctagt g 231

<210> 1215  
 <211> 284  
 <212> nucleic acid  
 <213> Zea mays

<400> 1215

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 ctgagtctca caagacccta acatctctga gcagtgtgac tgcttatggg tttgatcttg 180  
 tccgtggaac ccaaactctt gggcttgtca cgagtgtggt tttccctgct ggaaagtacc 240  
 tctttgctgg tgttgtggat ggacgcaaca tctgggctga tgat 284

1215  
 284  
 nucleic acid  
 Zea mays  
 1216  
 280  
 nucleic acid  
 Zea mays  
 1216  
 60  
 120  
 180  
 240  
 280

<210> 1216  
 <211> 280  
 <212> nucleic acid  
 <213> Zea mays

<400> 1216

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 gtctcttgag gctgttgttg ggaaggacaa gcttgtcgta tcaacttcct gctcgctcat 180  
 gcacaccgct gtggatcttg tgaacgagac taagcttgac agcgagatta agtcttggct 240  
 gcttttgcctg ccagaaagg tgttgagggt gatgctcttg 280

<210> 1217  
 <211> 283  
 <212> nucleic acid  
 <213> Zea mays

<400> 1217

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 tcatggactg gaggagagat tgggtttgac acctacttct ccatggccag gggcaacgcc 120  
 actgtccctg ctatggagat gaccaagtgg tttgacacca actaccactt tattgtccct 180  
 gaattgggcc caaacaccaa gttctcctat gcttctcaca aggctgttaa cgaatataag 240

gaggctagggc gctcgggtggt gatacgtgcc agtattgntg gac 283

<210> 1218  
 <211> 337  
 <212> nucleic acid  
 <213> Zea mays

<400> 1218

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 ancttgattc tgacaaattg gctgcattct ctgctgcata cgcagaactt gaatctgtac 180  
 tttctggatt gaatgtgctt gttgagactt actttgctga tgntcctgct gantcctaca 240  
 agaccctaac atctctgagc agtgtgactg cttatggttt tgatctgtcc gtngaaccca 300  
 aaccttgggc ttgtcacgag tgctggntcc ctgctgg 337

<210> 1219  
 <211> 446  
 <212> nucleic acid  
 <213> Zea mays

<400> 1219

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 agatcgcgga ccgcgtaaac aagatgctcg ccgtgctgga caccaacatc ctctgggtga 120  
 accctgactg cgggtctcaag acacgcaagt acacggaggt gaagcctgcc ctgaccaaca 180  
 tgggtctccgc caccaagctc atccgcaccc agctcggcag cgcgaaatga gtcagatttg 240  
 gtagatccat gggtttgttg acacaaaagc cagcagggtc atttgtttcg aataatttg 300  
 gtccttactc ctgttccatg gcgttaggtt aagcctcatt ggtgttggtg gggtagcggc 360  
 gttacgtgtc cgtgtttctaa agtttggaat gtgtgtttct ctttttttgt nggggtttgn 420  
 gtgccttttg gtatataaac agaaat 446

<210> 1220  
 <211> 228  
 <212> nucleic acid  
 <213> Zea mays

<400> 1220

ggcaaagaag atctccgagg aggaatacat cagtgccatc aangaagaaa tcagcaaggt 60  
 tgtcaagntc caagaggagc ttgacontga tgtgcttggt catggagagc cagaaagaan 120  
 tgacatgggt gagtacttcg gtgagcantt atctggtttt gcantcactg ccaacggntg 180  
 ggtgcnatct ttatggatca cgttgtgtga agccaccntt atctatgg 228

<210> 1221  
 <211> 232  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1221

caacttcctg ctgcgtcatg cacaccgctg tggatcttgt gaacgagact aagcttgaca 60  
 gcgagattaa gtcttggtt gcttttgctg cccagaaggt tgttgagggt gatgctcttg 120  
 ctaaggcatt ggctgggtcaa aaggatgagg cttacttcgc agcaaagtct gctgctcagg 180  
 cctcgaggaa atcctcacc cgtgtgacca atgaagaagt ccagaaggct gc 232

<210> 1222  
 <211> 217  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1222

aggatcttna ggctgntggt attcaggnga tccagattga tgaggcagct ctaaggnagg 60  
 gtctgccact gcgcaagtca gagcatgcat tctacctgga ctgggctgtc cactctttca 120  
 gaatcacnaa ctgcggagtc caggacacca cccagatcca naccatattg tgctactcca 180  
 acttcaacga catcatccac tccatcatcg acatgga 217

<210> 1223  
 <211> 277  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1223

ggcgctcggg gttgataccg tgccagtact tgttgagcca gtctcgtacc tgttgctctc 60  
 aaagcccgcc aagggtgtgg agaagggtt cctcttctt tcccttctta gcagcatcct 120  
 cccagtctac aaggagggtca ttgctgagtt gaaggcagct ggggcttcat ggattcagtt 180

tgatgagccc actcttgtcc tcgaccttga ttctgacaaa ttggctgcat tctctgctgc 240  
 atacgcagaa cttgaatctg tattntgant tnatggg 277

<210> 1224  
 <211> 206  
 <212> nucleic acid  
 <213> Zea mays

<400> 1224

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 aagatccaag aggagcttga cattgatgtg cttgtgcatg gagagccaga gagaaatgac 120  
 atgggttgagt acttcgggtga gcaattatct ggttttgcgt tcaactgccaa cggatgggtg 180  
 caatcctatg gntcacgctg tgtgac 206

<210> 1225  
 <211> 354  
 <212> nucleic acid  
 <213> Zea mays

<400> 1225

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 tgcagtctta tgggtcaagg tgngtcaagc ctgccgattc atctactgna acgtgtatgc 120  
 cgccaccaac nccatgaccg ttttctggtc gnnnacggcg cagagcatga cgtctcgccc 180  
 aatgaatggg atgntgactn gncagtcac antcctcaac tngtctctncg tcggaatgac 240  
 cagccgaggt ntganacttg ctaccagatc gctctngcga tnaagnnggn agtcgaggtc 300  
 tcgacgtggn gtatcaggtg accaantcga cnagctgcat gagagagagc gcgc 354

<210> 1226  
 <211> 228  
 <212> nucleic acid  
 <213> Zea mays

<400> 1226

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 cgtcaagcct ccgatcatct acggtgacgt gagccgcccc aaccccatga ccgttttctg 120  
 gtogaagagg cgcagagcat gacgtctcgc ccaatgaagg ggatgctgac tggcccagtc 180



acaatcctca actggtcctt cgtccggaat naccagccga ggttttat 228

<210> 1227  
<211> 260  
<212> nucleic acid  
<213> Zea mays

<400> 1227

cgnacntggg tntncgtcag gaacgaccag cctaggtttg agacatgcta ccaaatagct 60  
cttgcaatca aaaaggaggt tgaggatctt gangctgctg gtaantcagg tgatccagat 120  
cgatgaggca gctctaaggg agggctctgcc actacgcaag tcagagcatg cattctacct 180  
ggactgggct gtccactctt tcaggatcac caactgcgga gtccaggaca ccaccagat 240  
ccacaccac atgtgctact 260

<210> 1228  
<211> 276  
<212> nucleic acid  
<213> Zea mays

<400> 1228

cgtactatga ccaggtcctt gacaccaent gccatgctcg gcgctgtccc cgaacgctac 60  
tcatggactg gaggagagat tgggtttgac acctacttct ccattggccag gggcaacgcc 120  
actgtccctg ctatggagat gaccaagtgg ttgacacca actaccactt tattgtccct 180  
gaattgggcc caaacaccaa gttctoctat gcttctcana aggctgttaa ngaatataag 240  
aggctaaggc gctcgggtgtt aanaccggcc aatant 276

<210> 1229  
<211> 256  
<212> nucleic acid  
<213> Zea mays

<400> 1229

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ggattccctc ttctttccct tcttagcagc atcctcccag tctacaagga ggatcattgct 120  
gagttgaagg cagctggggc ttcattggatt cagtttgatg agcccactct tgtcctcgac 180  
cttgattctg acaaattggc tgcattctct gctgcatacg cagaacttga atctgtactt 240

tctggattga atgtgc

256

<210> 1230  
<211> 257  
<212> nucleic acid  
<213> Zea mays

<400> 1230

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ttctggctga agaggcacag agcatgacgt ctgcaccaat gaaggggatg ctgactggcc 120  
cagtcacaat cctcaactgg tccttcgtcc ggaatgacca gccgagggtt gagacttgct 180  
accagatcgc tcttgccgac aagaaggaag tcgaggatct tgaggctggt ggtattcagg 240  
ttatccaaat cgacgag 257

<210> 1231  
<211> 344  
<212> nucleic acid  
<213> Zea mays

<400> 1231

gggattccct cttctttccc ttcttagcag catcctccca gtctacaagg aggtcattgc 60  
tgagttgaag gcagctgggg cttcatggat tcagtttgat gagccactc ttgtcctcga 120  
ccttgattct gacaaattgg ctgcattctc tgctgcatac gcagaacttg aatctgtact 180  
ttctggattg aatgtgcttg ttgagactta ctttgctgat gttcctgctg agtcctacaa 240  
gaccctaaca tccctgagca gtgtgactgc ttatggtttg acttgctccgt ggaacccaaa 300  
ctcttggggt gtcacgagtg ctggtttccc tgctggaagt actc 344

<210> 1232  
<211> 278  
<212> nucleic acid  
<213> Zea mays

<400> 1232

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agtctttctg ggatgggaag annaccgctg aggatctgga gaaggttgct accgacctca 120  
gggccagtat ctggaagcag atggnngatg ctggggatca agtacatccc cagcaacact 180

ttctcgtact atgaccaggt ccttgacacc actgccatgc tcggcgctgt ccccgaaact 240  
tactcatgga ctggaggaga gattggattt gacactac 278

<210> 1233  
<211> 249  
<212> nucleic acid  
<213> Zea mays

<400> 1233

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agattaagtc ttggcttgct tttgctgccc agaaggttgt tgaggttgat gctcttgcta 120  
aggcattggc tgggtcaaaag gatgaggctt actcgcagca aatgctgcag ctcaggcctc 180  
gaggaaatcc tcaccccgctg tgaccaatga agaagtcag aaggctgctg ctgctctcaa 240  
gggctctga 249

<210> 1234  
<211> 295  
<212> nucleic acid  
<213> Zea mays

<400> 1234

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ccacgttcca aaggantgct ggcgtcccat attgttggat accntcgcac gggccccaag 120  
agagagctca agtttgctt ggagtcttcc tgggatggga agagcagcgc cgatgattta 180  
gagaaagttg cctctgacct caggtctagc atctggacng ccatgtcag angtgggnt 240  
cnagtacatt cccagcaaca ccttctcgta ctatgaccag gtctttgata ccagg 295

<210> 1235  
<211> 267  
<212> nucleic acid  
<213> Zea mays

<400> 1235

aacccaaact cttgggcttg tccacgagtg ctggtttccc tgctggaaaag tacctctttg 60  
ctgggtgttg ggatggacgc aacatctggg ctgatgatct tgctacatct ctcagcactc 120  
tccagtctct tgaggctgtt gttgggaagg acaagcttgt cgtatcaact tctgctcgc 180

tnntgcacac cgctgtggat cttgtgaacg agactaagct tgacagcgag attaagtctg 240  
gctgcttttg ctgcccagaa ggttggt 267

<210> 1236  
<211> 292  
<212> nucleic acid  
<213> Zea mays

<400> 1236

attggccctg gtgtctacga catccactct cctaggattc cctccaccga ggagatcgca 60  
gaccgcgtcg agaagatgct cgccgtgctg gacaccaaca tcctctgggt gaaccctgac 120  
tgtggtctca agacacgcaa gtacacggag gtcaagcccg cctgaccaa catggtctcg 180  
gccancaagc tcatccgcac ccagcttgcc aacgcgaaat gagtcgtttg atagctccat 240  
ggtttgatag cgcggaatga gccagttggt ttgaataatt tgggtgntac cc 292

<210> 1237  
<211> 217  
<212> nucleic acid  
<213> Zea mays

<400> 1237

ccaagaanan cttgacattg atgtgcttgt gcatggcgag cctnagagaa acgatatggt 60  
tgagtacttc ngtgagcagc tctccggttt tgcggttact gccaatggct ggggtgcagtc 120  
ttatgggtca aggtncgtca ngcctccgnt catctacggt gacgtgagcc gcccacacc 180  
catgaccgtn ttctggtcgn agactgcgca gagcatg 217

<210> 1238  
<211> 456  
<212> nucleic acid  
<213> Zea mays

<400> 1238

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ngaggntatc gccgaccgca tcgacaagat gctggcggtg ctggacacca acatcctctg 120  
gggtgaacccc gactgcggcc tcaagacccg caagtacacg gaggtcaagc ccgccctgac 180  
caacatggtc tccgcccgtc agtcatccn caccagctc gccagcgcca agtgagcacc 240

ttttttttgc ctttttctgt tccgaggagg gcgtcgtcna ntgccaattt ggttncaata 300  
 aacagggctn tgcccgccgt tctgtgttac tccgtctgtg gttagggttag tagttatott 360  
 gatctcgcn ncaacgccgt acctntttac tactctctgt ttggtgggtt tctgaagcaa 420  
 gttgcccntt tacttgtatn gtaaaaaccc aatngt 456

<210> 1239  
 <211> 289  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1239

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 ataccctcnc atgggccccca agagggagct caagtttgca cttgagtctt tctgggatgg 120  
 gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggcca gtatctggaa 180  
 gcagatggct gatgctggga tcaagtacat cccagcaac atttctcgta ctatgaccag 240  
 tcctgacacc atgccatgtc ggcgctgtcc ccgaaggtag tcatggatg 289

<210> 1240  
 <211> 274  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1240

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 gttgctctca aagcctgccca aggggtgtgga gaagggattc cctcttcttt cccttcttag 120  
 cagcatcctc ccagtctaca aggaggtcat tgctgagttg aaggcagctg gggcttcatg 180  
 gattcagttt gatgagccca ctcttgctct ngacctgaat tcggnacaaa tnggctgcat 240  
 tctcngcngc atangcngaa nttgaatctg tact 274

<210> 1241  
 <211> 268  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1241

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ggataccctc gcatgggncc cnaganggag ctcaagtttg cacttgagtc tttctgggat 120  
 ggganagagca acgctgagga tctggagaag gttgctaccg acntcagggc cagtatccgg 180  
 aagcanatgg cngatnctgg gatcaagtac atccccagca anacnttctc gnantatgac 240  
 caggtcttga naccatgccca tggctcgg 268

<210> 1242  
 <211> 260  
 <212> nucleic acid  
 <213> Zea mays

<400> 1242

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 tgctggcggt gctggacacc aacatcctct gggatgaacc cgactgcggc ctcaagacct 120  
 gcaagtacac ggaggtcaag cccgcctga ccaacatggt ctccgccgct aagctcatcc 180  
 gcaccagct cgccagcgcc aagtgagcac cttttttttg cttttttcgt ttccgaggag 240  
 ggcgtcgtcg atgccaattt 260

<210> 1243  
 <211> 280  
 <212> nucleic acid  
 <213> Zea mays

<400> 1243

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 gcatcgcccc cgggtgtctac gacatccact cccccaggat cccgtcggct gaggagatcg 120  
 ccgaccgcat cgacaagatg ctgngtgct ggacaccaac atcctctggg tgaaccccca 180  
 ctgcggcctc aagaccgca agtacacgga ggtcaagccc gccctgacca ncatggtctc 240  
 ngccggtaag ctnatccgca ccagctcgc naggccaagt 280

<210> 1244  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays

<400> 1244

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 ctgctgagtc ctacaagacc ctaacanctc tgagcagtgt gactgctnat ggttttgata 180  
 ttgtccgtgg aaccctaaact ctnggggctt gtcacgagtg ctgggtttcc ctgctggaaa 240  
 gtacctcttt gctggtgttg tggatggacn caacagctgg ggtngatgnt cttncnnacn 300  
 tcgcc 305

<210> 1245  
 <211> 198  
 <212> nucleic acid  
 <213> Zea mays

<400> 1245

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 atgangggga tgctgactgg cccagtcaca atcctcaact ggtccttcgt ccggaatgac 120  
 cagccgaggt ttgagacttg ctaccagatc gctcttgca tcaagaagga agtcgaggat 180  
 ctcgaggctg gtggtatt 198

<210> 1246  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays

<400> 1246

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 gcatacgag aacttgaatc tgtacttctg gattgaatgt gctgttgaga cttactttgc 180  
 tgatgttctt gctgagtcct acaagaccct aacatctctg agcagtgtga ctgcctatgg 240  
 ttttgatctt gtccgtggaa cccaaactct tgggcttgct acgagtgtg gtttccc 297

<210> 1247  
 <211> 286  
 <212> nucleic acid  
 <213> Zea mays

<400> 1247

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gagaagctgc tctccgtgtt ccgcgagggc gtcaagtacg gcgcgggcat cggccccggg 120  
 gtctacgaca tccactcnnc caggatccccg tcggctgagg agatcgccga ccgcatcgac 180  
 aagatgtgng gtgctggana cnaacatcct ctnggtgaac cccgactgog gcctcaagan 240  
 ccgcaagtac acngagggtca aancggngctg accaacangt ctccgt 286

<210> 1248  
 <211> 343  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1248

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 tgacgtctca gcccaatgaa nggggnntg actggccaag tcacaatncc tcaactggtc 120  
 cttcgtccgg aatgaccagc cgaggtttga gacttgctac cagatcgctc ttgcgatcaa 180  
 gaaggaagtc gaggatctcg aggctggtgg tattcaggtt atccaaatcg acgaggctgc 240  
 actgagagaa gggctgccgc tccgcaaggc tgnagcatgc atcttacttg gatgggctgt 300  
 tccattcctt cagaatcacc aatgcgaatc caggacacca ccc 343

<210> 1249  
 <211> 210  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1249

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 ggtattcagg tgatccagat cgatgaggca gctctaaggg agggctctgcc actacgcaag 120  
 tcagagcatg cattctacct ggactgggct gtccactctt tcaggatcac caactggcga 180  
 gtccaggaca ccaccagat ccacaccac 210

<210> 1250  
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 <213> Zea mays  
 <400> 1250

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aaggatgagg cttacttcgc agcaaagtct gcagctcagg cctcgaggaa atcctcacc 120  
 cgtgtgacca atgaagaagt ccagaaggct gctgctgctc tcaagggtc tgaccaccgc 180  
 cgtgcgacca acgttagtgc tagntt 206

<210> 1251  
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 <212> nucleic acid  
 <213> Zea mays

<400> 1251

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 ntgccgctcc gcaaggctga tcatgcattc tacttggact gggctgtcca ntcttcagn 180  
 attaccaact gcgagatcca ggacaccacc cagatccaca ccc 223

<210> 1252  
 <211> 186  
 <212> nucleic acid  
 <213> Zea mays

<400> 1252

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 ggaatacgtc actgccatca aggaggaaat caacaaggct gttaagcttc aagaagagct 120  
 tgacattgat gtgcttgtgc atggcgagcc tgagagaaac gatatggttg agtacttcgg 180  
 tgagca 186

<210> 1253  
 <211> 238  
 <212> nucleic acid  
 <213> Zea mays

<400> 1253

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 actgacctga ggtctagcat ctgnagcaa atgtcagang ctgggatcaa gtacattccc 180  
 aggcaatcac cttctcgtac tacgaccagg ttcttgatac cacggccatg cttggcgc 238

<210> 1254  
 <211> 277  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1254  
  
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 agctgctttg aggggatctg accaccgccc ttctaccact gtttctgcta gattggatgc 180  
 tcagcagaan aagctcaacc ttctgtcct tccacaacc acaattgggt cattccctca 240  
 gactgtggaa tcaggagggt tcgccgtgaa tacaagg 277

<210> 1255  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1255  
  
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 gtcgtcctct tccccggcg ctactctccc gctccacgtt ccaaaggaaa gatggcgctcc 120  
 catattgttg gataccctcg catgggcccc aagagagagc tcaagtttgc cttggagtct 180  
 ttctgggatg ggaagagcag cgccgatgat ttagagaaag ttgcctctga cctcaggctc 240  
 agcatctgga agcaaattgc agaagctggg atcaagtaca ttcccag 287

<210> 1256  
 <211> 219  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1256  
  
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 taaggcgctc ggtgttgata ccgtgccagt acttggttga ccagtctcgt acctgttgct 120  
 ctcaaagcct gccaaagggtg tggagaaggg attccctctt ctttcccttc ttagcagcat 180  
 cctcccagtc tacaaggagg tcattgctga gttgaaggc 219

<210> 1257

<211> 269  
 <212> nucleic acid  
 <213> Zea mays

<400> 1257

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 agagaagggc tgccgctccg caaggctgag catgcattct acttggaactg ggctgtccac 180  
 tccttcagaa tcaccaactg cggagatcca ggacaccacc catatccaaa cccanattgt 240  
 ctattcnanc ttnaacgaca tcatccact 269

<210> 1258  
 <211> 213  
 <212> nucleic acid  
 <213> Zea mays

<400> 1258

caacttcctg ctcgctcatg cacaccgctg tggatcttgt gaacgagact aagcttgaca 60  
 gcgagattaa gtcttggett gcttttgctg cccagaaggt tgttgagggt gatgctcttg 120  
 ctaaggcatt ggctgggtcaa aaggatgagg cttacttcgc agcaaattgt gctgctcagg 180  
 cctcgaggaa atcctcacc cgtgtgacca atg 213

<210> 1259  
 <211> 269  
 <212> nucleic acid  
 <213> Zea mays

<400> 1259

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 ctgcgatggg cccaagagg gagctcaagt ttgcacttga gtctttcttg gatgggaaga 180  
 gcaccgctga ggatctggag aaggttgcta ccgacctcag ggccagtatc tggaagcaga 240  
 tggctgatgc tgggatcaag tacatcccc 269

<210> 1260  
 <211> 266  
 <212> nucleic acid

<213> Zea mays

<400> 1260

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gaaggaagtt cgaggatctt naggtcgttg gtattcaggt tatccaaatc cgacgaggct 180  
gcactgagna gaagggtgc cgtccgcaa ggccgagcat gccatctact tgnactgggc 240  
tgtccactcc ttcagattac caactg 266

<210> 1261

<211> 443

<212> nucleic acid

<213> Zea mays

<400> 1261

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gaggagatcg ccgaccgcat cgacaagatg ctggcgggtgc tggacaccaa catcctctgg 120  
gtgaaccccg actgcggcct caagaccgc aagtacacgg aggtcaagcc cgcctgacca 180  
acatgggtctc cgcgctaag ctcatccgca ccagctcgc cagcgccaag tgagcacott 240  
atntttgcnt ntttngtttc cgaggagggc gtngtcgatg ccaattngng ttcaataaac 300  
agggctgtgc ccgcccgttc tgtngactnc gctgtgggtt aagttagtan tttcttngat 360  
ctcgnccca ccggtacct ggtttactac tctctgtttg gtggttcttg aagcaagtta 420  
cnnctgtac ttggatagaa aaa 443

<210> 1262

<211> 258

<212> nucleic acid

<213> Zea mays

<400> 1262

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gaatgtgctt gttgagactt actttgctga tgttctgct gagtcctaca agaccctaac 180  
atctctgagc agtgtgactg cttatggttt tgatcttgct cgtggaacct aaactcttgg 240

gcttgtcacg agtgctgg 258

<210> 1263  
 <211> 275  
 <212> nucleic acid  
 <213> Zea mays

<400> 1263

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 gaatgtgctt gttgagactt actttgctga tgttcctgct gagtcctaca agaccetaac 180  
 atctctgagc agtggtgactg cttatggttt tgatcttgct cgtggaaccc aaactcttgg 240  
 gcttgtcacg agtgctgggt tccctgctgg aaagt 275

gcttgtcacg agtgctgggt tccctgctgg aaagt

<210> 1264  
 <211> 253  
 <212> nucleic acid  
 <213> Zea mays

<400> 1264

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 gatgctcgcc gtgctcgaca ccaacatcct ctgggtgaac cctgactgtg gtctcaagac 120  
 acgcaagtac acggagggtca agcccgccct gaccaacatg gtctcggcca ccaagctcgc 180  
 ncgcacccag cttgccagcg cgaaatgagg tcgtttgata gctccatggt ctgatagcgc 240  
 ggaatgagcc agt 253

<210> 1265  
 <211> 211  
 <212> nucleic acid  
 <213> Zea mays

<400> 1265

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 tgcccagcag aagaagctta acctcccat ccttcccacc actacaatcg gttcgttccc 120  
 acaaaccgtt gagctcagga gggtcgccg tgagtacaag gcaaaaaaga tctctgagga 180  
 agaatacgtc actggccatc naggaggaaa t 211

<210> 1266  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1266  
  
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 gaccttgatt ctgacaaatt ggctgcattc tctgctgcat acgcagaact tgaatctgta 120  
 ctttctggat tgaatgtgct gttgagactt actttgotga tgttcttgct gagtccctaca 180  
 agaccctaac atctctgagc agtgtgactg cctatggttt tgatcttgct cgtggaaccc 240  
 aaactcttgg gcttgtcacg agtgctgggt tccctgctgg aaagtactct tgctggt 297

1266  
 297  
 nucleic acid  
 Zea mays  
 1267  
 267  
 nucleic acid  
 Zea mays  
 1267  
 60  
 120  
 180  
 240  
 267

<210> 1267  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1267  
  
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 gcttgtcacg agtgctgggt tccctgctgg aaagtacctc ttgctgggtg ttgtggatgg 120  
 acgcaacatc tgggctgatg atcttgctac atctctcagc actctccagt ctcttgaggc 180  
 tgttgttggg aaggacaagc ttgtcgtatc aacttctctg tcgctcatgc acaccgctgt 240  
 ggatcttggtg aacgagacta agcttga 267

<210> 1268  
 <211> 224  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1268  
  
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 tacnctcgca tggnccccaa nagggagctc aagtttgac ttnagtcttt cngggatngg 120  
 aagancacng ctgaggatcn cgagaaggtn gcnaccgacc tcagggccag tatctggaag 180  
 cagatngctg annctgggat caantacatn cccagcaaca cttt 224

<210> 1269

<211> 212  
<212> nucleic acid  
<213> Zea mays

<400> 1269

nnnnnnnnnn nnnnnngaca tggacgcgga cgtgatcacc atcgagaact cgcggtccga 60  
cgagaagctg ctctccgtgt tccgtgaggg tgtcaagtac ggcgcgggca tcggccccgg 120  
tgtctacgac atccaactccc ccaggatccc gtgggcccag gagatcgccg accgcatcga 180  
caagatgctg gccgtgctgg acaccaacat ca 212

<210> 1270  
<211> 292  
<212> nucleic acid  
<213> Zea mays

<400> 1270

cgagangctg ctctccgtgt tcngecnnggg cgtcaagtac ggngcgggcn tcggcctcgg 60  
tgtctacgac atennctccc ccaggntncc gtcnngctng gagatcgccg accgcatcga 120  
caagatgctg gcggtgctgg acaccaacat cctctgggtg aaccccgact cggcctcaag 180  
acccgcaagt acacggnggt caagcccgcc ctgaccaaca tggntnccgcc gctnagtcac 240  
ccgaccacgc tcggnagccg cgangtgagc acttttttgc cttttcgttc cg 292

<210> 1271  
<211> 287  
<212> nucleic acid  
<213> Zea mays

<400> 1271

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tggtgttggg aaggacaagc ttgtcgtatc aacttcctgc tcgctcatgc acaccgctgt 120  
ggatcttgtg aacgagacta agcttgacag cgagattaag tottggttg cttttgctgc 180  
ccagaagggt gttgaggtga tgctcntgct aaggcattgg ctgggtcaaaa ggatgaggnt 240  
acttcgcagc aaatgcngca gctcaggncn cgaggaaatc ctcaccc 287

<210> 1272  
<211> 265  
<212> nucleic acid

<213> Zea mays

<400> 1272

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ctccacgttc caaaggaaag atggcgtaen catattgntg gataccctcg catgggcccc 120  
aagagagagc tcaagtttgc cttggagtct ttctgggatg ggaagagcag cgccgatgat 180  
ttagagaaaag ttgcctctga cctcaggtct agcatctgga agcanatgtc agaagctggg 240  
atcaagtaca ttcccagcaa cacct 265

<210> 1273

<211> 223

<212> nucleic acid

<213> Zea mays

<400> 1273

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atggattcag tttgatgagc ccactcttgt cctcgacctt gattctgaca aattggctgc 120  
attctctgct gcatacgcag aacttgaatc tgtactttct ggattgaatg tgcttggtga 180  
gacttacttt gctgatgttc ctgctgagtc ctacaagacc cta 223

<210> 1274

<211> 261

<212> nucleic acid

<213> Zea mays

<400> 1274

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gtatctggaa gcagatggct gangctggga tcaagtacat cccagcaac actttctcgt 120  
actatgacca ggtccttgac accactgcca tgctcggcgc tgtccccgaa cgttactaca 180  
tggaactggag gagagattgg atttgacacc tacttctcca tggccagggg caagccactg 240  
ttctctgctat ggagatgacc a 261

<210> 1275

<211> 283

<212> nucleic acid

<213> Zea mays



<400> 1275

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ccaggctnnt gagngctggt gttgggaatg acaagottgt cgtatcaact tctgctcgc 120  
tcatgcacac cgctgtggat cttgtgaacg agactaagct tgacagcgag attaagtctt 180  
ggcttgcttt tgctgcccag aaggttgttg aggttgatgc nctgctaagg catggctggg 240  
caaaaggatg aggttactt cgcagcaaatt gctgcagctc agg 283

<210> 1276

<211> 185

<212> nucleic acid

<213> Zea mays

<400> 1276

gcgtcccaca ttgttggata cccnagcat gggcccgaag agggagctac aagtttgac 60  
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tcagggccag tatctggnag cagatggctg atgctgggat caagtacatc cccagcaaca 180  
ctttc 185

<210> 1277

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 1277

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tgctgcatac gcagaacttg aatctgtact ttctggattg aatgtgcttg ttgagactta 120  
ctttgctgat gttcctgctg agtcctacaa gaccctaaca tccctgagca gtgtgactgc 180  
ttatggtttt gatcttgtcc gtggaacca aactcttggg cttgtcacga gtgctgggtt 240  
ccctgctgga aagtactctt tgctgggtgt gtggatga 278

<210> 1278

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1278

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acaccaacat cctctgggtg aaccccgact ggggcctcaa gaccgcgaag tacacggagg 120  
tcaagcccg cctgacaaaa atgggtccccg ctganaagct cattcgcacc cagctcgcca 180  
gcgccaagtg agcacctttt tttgcctttt tcgtttccga ggagggcgtc gtcgatgcc 240  
tttgtttcca ataaacaggg cccccccct gtgcgc 276

<210> 1279  
<211> 267  
<212> nucleic acid  
<213> Zea mays

<400> 1279

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ttggctgcat tctctgtgc atacgcagaa cttgaatctg tactttctgg attgaatgtg 120  
cttggtgaga cttactttgc tgatgttccg gctgagtcct acaagaccct aacatccctg 180  
agcagtgtga ctgcttatgg ttttgatctt gtccgtggaa cccaaactct tgggcttgtc 240  
acgagtgtg gtttccctgc tggaaag 267

<210> 1280  
<211> 271  
<212> nucleic acid  
<213> Zea mays

<400> 1280

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gatgctgact ggcccagtca caatcctcaa ctggtcttcg tccggaatga ccagccgagg 120  
tttganactt gctaccagat cgctcttgcg atcaagaagg aagtcgagga tctcgaggct 180  
ggtggtatca gttatccaaa tcgacgaggc tgcagagag aagggtgccg ctccgcaagg 240  
ctgagcatgc attctattgg actgggtgtc a 271

<210> 1281  
<211> 188  
<212> nucleic acid  
<213> Zea mays

<400> 1281

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gctgatgata ttgctacata tctcagcaact ctccagtctc ttgaggctgt tgttggaag 120  
gacaagcttg tcgtatcaac ttctgctcg ctcatgcaca ccgctgtgga tcttgtgaac 180  
gagactaa 188

<210> 1282  
<211> 284  
<212> nucleic acid  
<213> Zea mays  
<400> 1282

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cacaatcctc aactggtcct tcgtccggaa tgaccagccg aggtttgaga cttgctacca 120  
gatcgctctt gcgatcaaga aggaagtcca ggatctcgag gctggtggta ttcaggttat 180  
ccaanttcga cgaggctgca ctgagagaag ggctgccgct ccgnaggntn agnatgattc 240  
naantgggan nggggcggcc nntcttnnag aatgnnaggn ggng 284

<210> 1283  
<211> 194  
<212> nucleic acid  
<213> Zea mays  
<400> 1283

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caaggcaaaa agatcaccca ggacgatacn tcagtgccat caaggaanaa tcagcaaggt 120  
tgtcaagatc caagaggagc ttgacattga tgtgcttggt catggagagc cagagagana 180  
tgacatgggt gagt 194

<210> 1284  
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<212> nucleic acid  
<213> Zea mays  
<400> 1284

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cgccaagggt gtggagaagg gattccctct tctttccctt cttagcagca tctcccagt 120

ctaaaggagg tcattgctga gttgaaggca gctggggctt catggattca gtttgatgag 180  
 cccatcttgt cctcgacctt gattctgaca aattggctgc attctctgct gcatacgag 240  
 aacttgaatc tgtatttctg gattgaatgt g 271

<210> 1285  
 <211> 209  
 <212> nucleic acid  
 <213> Zea mays

<400> 1285

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 cagtcacaat cctcaactgg tcttctgctc ggaatgacca gccgaggttt gagacttgta 180  
 ccagatcgct cttgctcaa gaaggaagt 209

<210> 1286  
 <211> 180  
 <212> nucleic acid  
 <213> Zea mays

<400> 1286

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 actggctcctt cgtccggaat gaccagccga ggtttgagac ttgctaccag atcgctcttg 120  
 cgatcaagac ggaantcgag gntctcnagg ctngtggtat tcaggttatc caaatcgagc 180

<210> 1287  
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 <212> nucleic acid  
 <213> Zea mays

<400> 1287

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 tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcagggccca 180  
 gtatctggaa gcagatggct 200

<210> 1288

<211> 324  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1288  
  
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 atctacggtg acgtgagccg ccccaacccc atgacgtttt ctggtcgaag acggcgcaga 180  
 gcatgacgtc tcgcncaatg aaggggatgc tgactggccc agtcacaatc ntcaactggt 240  
 cttcgtcggg acgaccagcc gaggtttgag attgctacca gatcgtctct gcgatnaaga 300  
 angaatcgan gatctgaagc tggc 324

<210> 1289  
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 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1289  
  
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 c 181

<210> 1290  
 <211> 275  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1290  
  
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 gaccaacatg gtctcggcca ccaagctcat ccgcacccag cttgccagcg cgaaatgagt 180  
 cgtttgatag ctccatggtc tgatagcgcg gaatgagcca gtttgttttg aataatttgg 240  
 gtgttaccce ctgttccatg gtgtnagtgt taggt 275

<210> 1291

<211> 202  
 <212> nucleic acid  
 <213> Zea mays

<400> 1291

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 tatggttttt atcttgtccg tggaacccaa actcttgggc ttgtcacgag tgctggtttc 180  
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 <213> Zea mays

<400> 1292

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<210> 1293  
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 <212> nucleic acid  
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<400> 1293

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 tttattgtcc ctgaacttgg cccaagcacc aagttcacat acgctnctca caaggctggt 180  
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 gc 242

<210> 1294  
 <211> 176  
 <212> nucleic acid  
 <213> Zea mays

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tcttgcgatg aagaaggaag tcgaggatct cgaggctggt ggtattcagg ttatcc 176

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actcttgctc tcgaccttga ttctgacaaa ttggctgcat tctctgctgc atacgcagaa 180  
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<210> 1296  
<211> 312  
<212> nucleic acid  
<213> Zea mays

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gttgataacc ctgcgatggg cccaagagg gagctcaagt ttgccttga gtctttctgg 180  
gatgggaaga gcagcgccga ggatttgag aaagttgcca ctgacctgag gtctagcatc 240  
tggaagcaaa tgtcagaagt gggatcaagt aattcccagc aatacttctc gtatacgncc 300  
aggttttgat ac 312

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<212> nucleic acid  
<213> Zea mays

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agcccactct tgtcctcgac cttgaatnct gaacaaattg gctgcattct ctgctggcct 180  
acgcagaact ngaatctgta ctntctggat tgaatgtgct tgttgagact tactttgctg 240  
atgttcctgc tgagtcctac aagaccctaa 270

<210> 1298  
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<212> nucleic acid  
<213> Zea mays  
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tgaaccccgga ctgcggcctc aagaccgcga agtacaggag g 161

<210> 1299  
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ctcaggcctc gaggaaatcc tcaccccgctg tgaccaatga agaagtccag aaggctgctg 180  
ctgtctcaag ggctctgacc accgccgtgg gac 213

<210> 1300  
<211> 194  
<212> nucleic acid  
<213> Zea mays  
<400> 1300

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tctgggatgg gaagagcacc gctgaggatc tggagaaggt tgctaccgac ctcaggggcca 180  
gtatctggaa gaaa 194



<210> 1301  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 gtcacgagtg ctggtttccc tgctggaaag tacctctnng ctggtgttgt ggatggacgc 120  
 aacatctggg ctgatgatct tgctacatct ctcagcactc tccagtctct tgaggctgtt 180  
 gttgggaagg acaagcttgt cgtatcaact tctgtctgc tcatgcacac cgctgtggga 240  
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<210> 1302  
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 <213> Zea mays  
  
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 caagcctccg atcatctacg gtgacgtgag cc 152

<210> 1303  
 <211> 152  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1303  
  
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 ncccacatgt gctactccaa cttcaacgac atcatccact cgatcatcga catggacgcg 120  
 gacgtgatca ccatngagaa ctcgngggnc gg 152

<210> 1304  
 <211> 468  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1304

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 gcatcctccc antctacaag gaggtcattg ctganttnaa ngcanctggg gcttnatgga 180  
 ttcagtttga tgancccaact cttgtcctcg accttgattc tgacaaattg gctgcattct 240  
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 ttactttgct tatgttcctg ctgaantcct acaaaaaccc taacatccct taaacanngt 360  
 gaactgctta ttgggttttta atcttttccc gnggaaccca aaactnttgg gcttntcacn 420  
 aatgctttgg ttttccttgc ttgaaaaaan cnnnctttgt tgggggtgg 468

<210> 1305  
 <211> 292  
 <212> nucleic acid  
 <213> Zea mays

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 gctcatgcac accgctgtgg atcttgtgaa cgagactaag cttgacagcg agattaagtc 180  
 ttggetgctt ttgctgcccc gaaggttgtt gaggttgatg ctctcgctaa ggcattggct 240  
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<210> 1306  
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 <212> nucleic acid  
 <213> Zea mays

<400> 1306  
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 gtgaaccccg actgnggnct naaaaacccg caagtacacg gaggtcaaag ccccgncctg 180  
 accaacaatgg tcttccgncc gctaagctca tccgcaccca gctngccagc gccaaagtga 240  
 cacctttttt ttgccttttt cgtttccgag gaagggcgtn gtcgatgcc aatttggtttc 300  
 aataaacang gctgtgcccc ccggtctggt gnactccgct gngggtaagg taagaagttt 360

cttganccttg ncccangccg gnactgggta ctactntntg gtngnggttc tgag 414

<210> 1307  
<211> 163  
<212> nucleic acid  
<213> Zea mays

<400> 1307

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tggataccct cgcattgggc ccaagagaga gctcaagttt gccttggagt ctttctggga 120

tgggaagagc agcgccgatg atttagagaa agntgnntcg gac 163

<210> 1308  
<211> 233  
<212> nucleic acid  
<213> Zea mays

<400> 1308

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cctcgtcccc cggcgctact ccccgctcc acgggtccaag gaaagatggc gtcccatatt 120

gttgataacc ctgcattggg cccaagagg gagtcaagt ttgccttga gtctttctgg 180

gatgggaaga gcagcgccga ggatttggag aaagttgcca ctgacctgag gtc 233

<210> 1309  
<211> 225  
<212> nucleic acid  
<213> Zea mays

<400> 1309

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ctcctcgtcc cccggcgcta ctccccgct ccacggcca aggaaagatg gcgtccata 120

ttgttgata cctcgcattg ggcccaaga gggagctcaa gtttgccttg gactctttct 180

gggatgggaa gagcagcgc gaggatttgg agaaagttgc cactg 225

<210> 1310  
<211> 303  
<212> nucleic acid  
<213> Zea mays

<400> 1310

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atattgttgg ataccctcgc atgggcccc agagggagct caagtttgcc ttggagtctt 180  
tctgggatgg gaagagcagc gccgaggatt ggagaaagtt gccatgacct gangttancc 240  
attggaagca aatgtcagaa ntgggatcaa gtacatccca gaatacttct cgtatagaca 300  
ggt 303

<210> 1311

<211> 293

<212> nucleic acid

<213> Zea mays

<400> 1311

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aggcagctgg ggcttcatgg attcagtttg atgagccac tcttgteccg gaccttgatt 180  
ctgacaaatt ggctgcattc tctgctgcat acgcagaact tgaatctgta ctttctggat 240  
tgaatgtgct gttgagacta cttgctgatg ttctgctga gtcctacaag acc 293

<210> 1312

<211> 311

<212> nucleic acid

<213> Zea mays

<400> 1312

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cgaggtttga gacttgctac cagatcgtct tgcgatcaag aaggaagtcg aggatcttga 180  
ggttgtgtat tcaggttatc caaatcgaga gctgcatgag agaggggtgc gctccgcaag 240  
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ngntcacanc c 311

<210> 1313

<211> 121  
 <212> nucleic acid  
 <213> Zea mays

<400> 1313

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 a 121

<210> 1314  
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 gttggatacc ctgcgatggg cccaagagg gagctcaagt ttgccttgga gtctttcttg 180  
 gaggggaaga gcagtgccga ggattnggag aaattgcnac tg 222

<210> 1315  
 <211> 190  
 <212> nucleic acid  
 <213> Zea mays

<400> 1315

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 gaccctaaca ncnngagca gtgtgactgc ntatggtttt gatcttgtcc gtggaacca 180  
 nactctnggg 190

<210> 1316  
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 <212> nucleic acid  
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<400> 1316

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gtctttctgg gatgggaaga acaccgctga agatctggag aaggttgcta ccgacctcag 120

<210> 1317  
<211> 159  
<212> nucleic acid  
<213> Zea mays

<400> 1317

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aatgnagaag tccaaaatgc tgengntgen ctcaanggct atgancancg ccgngggacc 120

aacgttagtg ctagattgga tgnccagcag aagaagctt 159

<210> 1318  
<211> 288  
<212> nucleic acid  
<213> Zea mays

<400> 1318

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ggccaagccc gccctgacca acatggtctc ggccaccaag ctcatccgca ccagcttgcc 120

cagcgcgaaa tgaggtcggt tgatagctcc atggtctgat agcgcggaat gagccagttg 180

ttttgaataa tttgggtggt accccctggt ccatggtggt agtggttaggt tagcctctca 240

ttggtgggat acgccgttac aagatgtggt ctaagtttgg agtgtgtg 288

<210> 1319  
<211> 268  
<212> nucleic acid  
<213> Zea mays

<400> 1319

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tgctacgggt acgttagccg ccnaacccc atgaccgttt tctcggtcga agacggcaca 180

gagcatgacc gtctcgccca atgaagggga tgctgactgg ccagtcaca atcctncaac 240

tggtcccttn cgtncgaaa tganccac 268

<210> 1320

<211> 118  
 <212> nucleic acid  
 <213> Zea mays

<400> 1320

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<210> 1321  
 <211> 151  
 <212> nucleic acid  
 <213> Zea mays

<400> 1321

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 ccaggttctt gataccacgg ccattgcttg cgtgttccca gagcgctact cttggactgg 120  
 aggtgagatt ggctgagcac tacttctcta t 151

<210> 1322  
 <211> 436  
 <212> nucleic acid  
 <213> Zea mays

<400> 1322

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 gcaagtcacg gaggtcaagc ccgccctgac caacatggtc tnggccacca agcttattcc 180  
 gcacccaact tgnacgcgcc gaaatgaagt cgtttgatag ctcatggtct gatagcgccg 240  
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 aaggtagcct ntnattggtg ggataccccc gttccagant nggtctaagg ttggantggg 360  
 tggttntctt tnggctatgg ttcttggggg attgtgtgtt ccttgggtat taacnggaat 420  
 tgaataccca gtcttt 436

<210> 1323  
 <211> 118  
 <212> nucleic acid  
 <213> Zea mays

<400> 1323

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tcaagtagcn cncgggcacn ngcacccggtt tctacnacat ccaactcccc angatccc 118

<210> 1324

<211> 324

<212> nucleic acid

<213> Zea mays

<400> 1324

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ggacacaaca tctctgggtg aaccccgact gcggctcaag accgcaatac acgagtcaag 180  
cccgctgac caacatggtc tccgcccgtc agtcatccgc accagtcgcc agcgccaagt 240  
gagcactttt tttgcttttc gttcgaggag ggtgtgatgc aaattgttcn ataaaanggt 300  
gtgccgccgt tctgttgat cgtt 324

<210> 1325

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 1325

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gcaccttttt tttgcctttt tcgtttccga ggagggcgtc gtcgatgcca atttgtttcc 180  
aataaacagg gctgtgccgc cgttctgttg tactccgtct gtggttaggt tagtagtttt 240  
cttgatctcg cccccacgcg gtacctgttt tactactctc tgtttg 287

<210> 1326

<211> 131

<212> nucleic acid

<213> Zea mays

<400> 1326

gtatgccnag cagaagaagc ttaaccnccc catccttccc accanngaca atcggttcgt 60



tcccanaaan cgttgagctc caggaggggc cgccgtgagt ncaaggcaaa aaagatctct 120  
gaggaggaat a 131

<210> 1327  
<211> 155  
<212> nucleic acid  
<213> Zea mays  
  
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tacttcggtg agcagctcct ccggttttgc gttcactgcc aatggctggg tgnagtctta 120  
tgggtcaang tgcgtcaagc ctccgatcat ctacg 155

1328  
264  
nucleic acid  
Zea mays  
1328

<210> 1328  
<211> 264  
<212> nucleic acid  
<213> Zea mays  
  
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tgcacttgag tctttctggg atgggaagag caccgtgagg attggagaag gttgtacgac 180  
ctcaggcagt attggaacag atgntgatnt ggataagtac atcccacaac actttgtnta 240  
tacagtcttg cacatgcant gggt 264

<210> 1329  
<211> 270  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1329

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agcagcatcc gccagtgt anaaggaggt cattgctgag ttgaaggcag ctggnggctt 180  
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tgctttntct gctgcatacg agaantggaa 270

<210> 1330  
 <211> 268  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1330  
  
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 tcttgtcctc gaccttgatt ctgacaaatt ggectgcatt ctctgctgca tacgcagaac 180  
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 tgantctaca agaccntnta tccntggg 268

<210> 1331  
 <211> 291  
 <212> nucleic acid  
 <213> Zea mays  
  
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 ttttttttgc ctttttcgtt tccgaggagg ggtcgtcgat gccaatgtgt ttccaataaa 180  
 cagggtgtg nccgccgttc tgttgtactc cgtctgtggt taggttagta nttttcatga 240  
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<210> 1332  
 <211> 128  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1332  
  
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 cttgagtc 128

<210> 1333  
 <211> 286  
 <212> nucleic acid

<213> Zea mays

<400> 1333

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tctgggatgg gnagagnacc gctgnaggat ctgganaagg ttgctaccga cntcaagggc 180  
cagtancctng angccngatg ggntgatgct gggatcaagt acatccncca gcaacaacct 240  
tgctctactt atgacaaggt ccttgacaca ctgccatgct cgcggt 286

<210> 1334

<211> 156

<212> nucleic acid

<213> Zea mays

<400> 1334

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acgcaagtac acggagggtca agcccgcctt gaacaa 156

<210> 1335

<211> 185

<212> nucleic acid

<213> Zea mays

<400> 1335

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aaggctgctg ctgctctcaa gggctctgac caccgctgtg naaccaacgt tatgctagat 180  
tggat 185

<210> 1336

<211> 139

<212> nucleic acid

<213> Zea mays

<400> 1336

ggagggtgag gatcttgagg ctgctggtat tcagggtgatc cagnttgatg aggcagctct 60

aaggganngt ngccactgcg caagtcccag agcatgcatt ctacctggac tgnngctgtcc 120  
actcttttcag antcaccaa 139

<210> 1337  
<211> 114  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1337

cgaggatctt gaggctgggtg gtattcaggt tatccaaatc gacgaggctg cactgagaga 60  
agggctgccg ctccgcaagg ctgaacatgc atctaactgg actgggctgt ncca 114

<210> 1338  
<211> 254  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1338

acggagggtca agcccgcctt gaccaacatg gtctccgctg ctaagctcat tcgcacccag 60  
ctgccagcgc ccaagtgagc accttttttt gccttttttcg tttccgagga gggcgctcgtc 120  
gatgccaaatt tgttttccaat aaacaggggt cccccctgt gcccgccgtt ctgtttgtgt 180  
cgtctgttg ttaggttact agttttcttg atctcgcccc caccggttac ctgttttact 240  
atctgttttg tggt 254

<210> 1339  
<211> 289  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1339

cggagggtcaa gcccgccctg accaacaatg tctccgccgc taagctcatc cgcacccagc 60  
tcgccagcgc caagtgagca cctttttttt gccttttttcg tttccgagga gggcgctcgtc 120  
gatgccaaatt tgttttccaat aaacaggggt gtgccgccgt tctgtttgtac tccgtctgtg 180  
gttaggttag tagttttctt gatctcgccc ccacgcggtta cctgtttttac tactctctgt 240  
ttggtgggtt ctgaggaagt tgcccgtgta ttgtatcgta aaaaccgag 289

<210> 1340

<211> 252  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1340  
  
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 ggtctccgcn gctaagctca tcngeacnea gctcgccagc gccaaagtng cacctttttt 120  
 ttgccttttt cgtttccgan gagggentcg tngatgccaa tttgtttcna ntaancaggg 180  
 ntgtncacgc cgttctgttg nactcgtct gtggttagg tagtagtttt cttgatctcg 240  
 cccccaagcg nt 252

<210> 1341  
 <211> 79  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1341  
  
 ctgcggagtc caggacacca ccagatcca naccacatg tgtacncaa cttcaacgac 60  
 ntctcnant ccatnanng 79

<210> 1342  
 <211> 260  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1342  
  
 cggagggtcaa gcccgccctg accaacaatgg tctccgctgn taagctcatt cgcaccacgc 60  
 tcgncagcgc caagtgaaga cctttttttg cctttttcgt ttccgaggag ggcgctgctg 120  
 atgccaattt gtttccaata aacagggctc cccccctgtg cccgcngttc tgttgtgtgc 180  
 cgtctgtggt taggttacta gttttcttga tctcgcccc acgcggtacc tgttttacta 240  
 tctgtttggt gggtttctgag 260

<210> 1343  
 <211> 124  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1343

caagggtctc gcaccaccgc cgtgcgacca acgttagtgc tagattggat gccagcaga 60  
 agangttaac ctccccatcc ttcccaccan tacaatcggg ttcgttccca caaacggtt 120  
 agct 124

<210> 1344  
 <211> 505  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1344

agnttggttn agggggggng atnnaaagge natctcgtac cgggccggaa ttcccgggtc 60  
 gaccacgcgc tccggtgaa ctagagtcgg cattctctgg attgaatgtg cttattgaga 120  
 catactntgc ngacattcct gctgagtcct acaagaccct cacgtcattg agtggtgtga 180  
 ctgcttacgg ttttgatott atcgtggag ccaagaccct tgatcttattc aggagcagct 240  
 tcccctctgg gaagtacctc ttcgctgggtg ttgtggatgg acgcaacatt tgggctgatg 300  
 atcttgctgc atctcttagc actcttcagt ctcttgaggc cgttgctggc aaggacaaac 360  
 ttgtgggtgc aacctcctgc tcaactgatgc acaccgccgt gatcttgtaa atgagactaa 420  
 actggatgaa tgaagattaa gtcattggctt gcatttgctg cccaaaagggt tgtcgaggtt 480  
 aatgcccttg cgaagctttg gcaag 505

<210> 1345  
 <211> 300  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1345

aaacttggtg tgtcaacntc ctgctcactg atgcanaccg ctgttgacct tgtaaattgag 60  
 actaagctgg atgatgagat taagtcatgg cttgcatttg ctgccccaaa ggttggtgag 120  
 gttaatgcc ttgccaaggc nttggcaggc caaaaggatg aggtctactt tgcagccaat 180  
 gctgctgctc aggcctcaag gagatcatcg cccagggtga caaacganga ggtccagaag 240  
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<210> 1346  
 <211> 290  
 <212> nucleic acid

<213> Zea mays

<400> 1346

ggacgcaaca tttgggctga tgatcttgct gcatctctta gcactcttca ttctcttgag 60  
gctgttgctg gcaaggacaa acttggtgtg tcaacctcct gctcactgat gcacaccgct 120  
gttgaccttg taaatgagac taagctggat gatgagatta agtcatggct tgcatttgct 180  
gccccaaaagg ttgttgaggt taatgccctt gccaaaggctt tggcaggcca aaaggatgag 240  
gtctactttg cagccaatgc tgctgctcag gctcaagga gatcatcgcc 290

<210> 1347

<211> 259

<212> nucleic acid

<213> Zea mays

<400> 1347

aacctcctgc tcaatgatgc acaacgctgt tgaccttgta aatgagacta agctggatga 60  
tgagattaag tcaatggcttg catttgctgc ccaaaagggtt gttgagggtta atgcccttgc 120  
caaggctttg gcaggccaaa aggatgaggt ctactttgca gccaatgctg ctgtcaggc 180  
ctcaaggaga tcatcgccca gggtgacaaa cgaggaggctc cagaaggctg cagctgcttt 240  
gaggggatct gaccaccgc 259

<210> 1348

<211> 296

<212> nucleic acid

<213> Zea mays

<400> 1348

tgcttatcga gacatacttc gctgatattc ctgctgagtc ctacaagacc ctcacatcat 60  
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tcaggagcag cttccctctt gggaagtacc tcttcgctgg tgttgtagat ggacgcaaca 180  
tttgggctga tgatcttgct gcatctctta gcactcttca ttctcttgan gctgttgctg 240  
gcaaggacaa acttggtgtg tcaacctcct gctcactgat gcacaccgct gttgac 296

<210> 1349

<211> 272

<212> nucleic acid

<213> Zea mays

<400> 1349

cctgctgagt cctacaagac cctcaagtc ttagtggtg tgactgctta cggttttgat 60  
cttatccgtg gagccaagac ccttgatctt atcaggagca gcttcccctc tgggaagtac 120  
ctcttcgctg gtggttgga tggacgcaac atttgggctg atgatcttgc tgcattctt 180  
agcactcttc agtctcttga ggccgttgc ggcaaggaca aacttggtgt gtcaacctcc 240  
tgctcactga tgcacaccgc tgttgatctt gt 272

<210> 1350

<211> 265

<212> nucleic acid

<213> Zea mays

<400> 1350

acgtcattga gtggtgtgac tgcttacggt tttgatctta tccgtggagc caagaccctt 60  
gatcttatca ggagcagctt cccctctggg aagtacctct tcgctgggtgt tgtggatgga 120  
cgcaacattt gggctgatga tcttgctgca tctcttagca ctcttcagtc tcttgaggcc 180  
gttgctggca aggacaaact tgtggtgtca acctcctgct cactgatgca caccgctgtt 240  
gatcttgca atgaaactaa gctgg 265

<210> 1351

<211> 238

<212> nucleic acid

<213> Zea mays

<400> 1351

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cttgaggctg ttgctggcaa ggacaaactt gtggtgtcaa cctcctgctc actgatgcac 120  
accgctgttg accttgtaaa tgagactaag ctggatgatg agattaagtc atggcttgca 180  
tttgctgccc aaaaggttgt tgaggttaat gcccttgcca aggctttggc aggccaaa 238

<210> 1352

<211> 226

<212> nucleic acid

<213> Zea mays



<400> 1352

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cttgtgggtg caacctcctg ctcactgatg cacaccgctg ttgatcttgt caatgaaact 120  
aagctggatg atgagattaa gtcattggctt gcatttgctg cccaaaaggt tgcgaggtt 180  
aatgcccttg cgaaggcttt ggcaggccaa aaggatgagg cttact 226

<210> 1353

<211> 248

<212> nucleic acid

<213> Zea mays

<400> 1353

cctgctgagt cctacaagac cctcaantca gggantngtg tgactgctta cggttttgat 60  
cttatccgtg gagccaagac ccttgatctt atcaggagca gcttccctc tgggaagtac 120  
ctcttcgctg gtgttggtga tggacgcaac atttgggctg atgatcttgc tgcactctt 180  
agcactcttc agtctcttga ggcggttgct ggcaaggaca aacttggtgt gtcaacctcc 240  
tgctcatg 248

<210> 1354

<211> 203

<212> nucleic acid

<213> Zea mays

<400> 1354

cggacgnittg gcgcgagtgc tccctcctct gcngccatgg cgctctctgt ggagangacc 60  
tcgtctggan cngagtacaa gttcaaggtc tctcncangc ggacttcggc cgctcnaga 120  
ttgagctncc cnagttcgaa atgcccgncc tcatgnetnc nccncngatt ncgcnccttn 180  
aaancctnnn naggcgaana ttc 203

<210> 1355

<211> 96

<212> nucleic acid

<213> Zea mays

<400> 1355

ttgggcttna aaactnacct ngnttaagc gcattaccat taaaccccaa acnnancgt 60

tggtgtttcc ccagacnaaa actgngatan anntgc

96

<210> 1356  
<211> 518  
<212> nucleic acid  
<213> Zea mays

<400> 1356

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cccacgcgtc cgcnnacgcg tncgctnacn cgtcgcgaca ngcgtcngcn canncgttcg 120  
cccacnacgg negatnnttc natttccna tctcgtcana ttcaattcgc cgagttcttc 180  
ctctctctgc tcgncatggc gctctatgtg ganaanacct cgtcatggac gggagtacaa 240  
ggtcaaggat ctatagcang cngacttagn tcgccttgan attgatctgg ccnaggncga 300  
aatgcncgna ctcatnggnt gccncgcca gtttgcccg ncaagccct ttggcgntgc 360  
taggatctng gnntctcttt acatgaccat tnacacnggn gaagtgattg agacnnttac 420  
cgngctcgtt gccgaggtnc ngtggtgatc ntcaacatc ttattnacac angaccacgc 480  
ttnaancggn attggtncgc ganttggtg canaaatt 518

<210> 1357  
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<212> nucleic acid  
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<400> 1357

cggcaagatc cccgaccgg agtccaccga caacgctgag ttcaagatcg tgctcacnat 60  
catccgcgac gggctcaagg ctgaccccaa gaagtaccgc aagatgaagg agaggcttgt 120  
cggcgtctct gaggagacca ccacgggtgt caagaggctc taccagatgc aggagaccgg 180  
cgccctcttc ttccctgcca ttaacgtcaa cgattccgtc accaagagca agtttgacaa 240  
cctgtatggt tgccgccact cgctccctga tggctctgatg agggccactg acgttatgat 300  
cgccggaaaag gttgccgtgg tctgcggata cggtgatgtc ggcaagggtt gtgctgctgc 360  
ctcaagcagg ctggtgcccg tgcattgtg accgagatcg acccatctg tgcccttcaa 420  
gctctgatgg anggtcttta ngtcctt 447

<210> 1358

<211> 515  
 <212> nucleic acid  
 <213> Zea mays

<400> 1358

gnnnagnnnn nnnannnttt gnangggggg gggaagagan attttancgg tccggattcc 60  
 cgggtcgacc cacgcgtccg ccatcatccg cgacgggctc aaggctgacc ccaagaagta 120  
 ccgcaagatg aaggagaggc ttgtcggcgt ctctgaggag accaccacgg gtgtcaagag 180  
 gctctaccag atgcaggaga cgggcgcctt cctcttcctt gccattaacg tcaacgattc 240  
 cgtcaccaag agcaagtttg acaacctgta tggttgccgc cactcgctcc ctgatgggtc 300  
 gatgagggcc actgacgtta tgatcgccg taagggttgc gtgggtctgc gatacgggtga 360  
 tgtcggcaag ggttgtgtg ctgccctcaa gcaggctggt gcccggtgtca ttgtgaccga 420  
 gatcgacccc atctgtgccc tccaggctct gatggaaggg tcttcagggtc cttcccttgg 480  
 aagacgttgt ctctgaactg acatcttctg gacca 515

<210> 1359  
 <211> 530  
 <212> nucleic acid  
 <213> Zea mays

<400> 1359

ggnnnagggt gntttngtcc naanntntnt ttnncttctg ccngtaccgg tccggattcc 60  
 cgggtcgacc cacgcgtccg cccacgcgtc cgtgggtgac cacatgagga agatgaagaa 120  
 caatgccatt gtctgcaaca ttggccactt tgacaatgaa attgatatgc tcggccttga 180  
 gacctaccct ggcgtcaagc gcatcaccat caagccccag actgaccgct ggggtgtccc 240  
 cgagaccaac actggcatca ttgtccttgc tgagggtcgc ctgatgaacc ttgggtgtgc 300  
 tactggccat cctagctttg tcatgtcctg ctcatcact aaccagggtca ttgcccact 360  
 tgaactgtgg aaggagaaga gctctggcaa gtatgagaag aagggtgatg tgctcccaa 420  
 gcaccttgat gagaagggtg ctgctctcca cttgggcaag cttggtgcca agctgaccaa 480  
 gctcaccaag tctnagccga ctacatnanc gtgcccacgc agggtcctac 530

<210> 1360  
 <211> 501  
 <212> nucleic acid

<213> Zea mays

<400> 1360

ggnngnnngn aattctaacn tncngcagcc tgcanaagtc canggtccac ccacncgtcg 60  
ncgaggagga gtacgagaag accggcaaga tccccgaccc ggagtccacc gacaacgcga 120  
gttcaagatc gtgtcacca tcatccgga cgggctcaag gctgaccca agaagtacgc 180  
aagatgaagg agaggcttgt cggcgtctct gaggagacca ccacgggtgt caagagctct 240  
accagatgca ggagaccggc gccctcctct tccctgccat taacgtcaac gattcgtcac 300  
caagagcaag tttgacaacc tgtatgggtg ccgccactca ctccctgatg gtctatgagg 360  
gccaccgacg ttatgatcgc cggtaagggt gccgtggctt gcggatacgg tgagttggca 420  
agggttgtgc cgctgcactc aaancangct gggtgcccgt gtcaantgtg acgaagattt 480  
aaccaatct gcgncctcca a 501

<210> 1361

<211> 473

<212> nucleic acid

<213> Zea mays

<400> 1361

cgccgnaag gttgtgtgg tctgcggata cggatgatgc ggcaaggggt gtgtgtgtgc 60  
cctcaagcag gctggngccc gtgtcattgt gaccgagatc gaccccatct gtgccctcca 120  
ngctctgatg gagggctctc angtccttcc cttggaggac gttgtctctg aagctgacat 180  
cttcgtgacc accactggca acaaggatat catnatggat gaccacatga ggaagatgaa 240  
gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcgggct 300  
tnanacctac cctgncgtca agcgcacnac catcaagccc cagactgacn cgctggntgg 360  
nccccgagac caacactggg catcattgtc cttgctgaag gtcggctgat naaccttgng 420  
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<210> 1362

<211> 524

<212> nucleic acid

<213> Zea mays

<400> 1362

gnnnnnngggn ggngaattgtc tnccttngtcg tgcgggtcca gcgatacagg ggccccggac 60  
 ncgtccggga ncnatntccc catctcccag atccaattcg cgagttctcc ctctcttgcc 120  
 gccatggcgc tctcngtgga gaagacctcg tctggacggg agtacanggt caaggatctc 180  
 tcgcaggcgg acttcggccg cctcgagatt gagctggccg aggtcgaaat gcccggcctc 240  
 atggcgtgcc gcgccgagtt cggcccgctc aagcccttcg ccggcgctag gatctcgggg 300  
 ttctctccaca tgaccatcca ngaccggcgt cctcatcgag accctcaccg cgctcggcgc 360  
 cgangtccgc tgggtgctct gcaacatctt ctccacgcag gaccacgccg ccgccgccat 420  
 cgcgcgcgac tcggccgccg tgttcgctg gaanggggaa accctcgang agtactggtg 480  
 gtgcancgaa cgctgcctcg actggggggc angcgggccg cccc 524

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<210> 1363  
 <211> 478  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1363  
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 tccatttccc catctcccag atccaattcg cgagttctcc ctctcttgcc gccatggcct 120  
 ctctgtggag aagacctcgt ctggacggga gtacaaggtc aaggatctct cgcaggcgac 180  
 ttcgccgcc tcgagattga gctggccgag gtcgaaatgc ccggcctcat ggctgctgcg 240  
 ccgagttcgg cccgtogaag ccttcgctg gcgctaggat ctcggggtct ctccaatgac 300  
 catccagacc gccgtctca tcgagaccct caccgcgctc ggccgccgang tccgtggtgc 360  
 tcctgcaaca tttctccac gcaggaccac gccgccgcc ccctcgcgcg cgatcggccg 420  
 ccgtgttcgc ctggaagggg gaagacctcg aggagtactg gtggtgcacc gacgctgc 478

<210> 1364  
 <211> 528  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1364  
 gnnngnnnng nnnnnnnaag gtnntgttn ngnaaggagt tttatggnat actcgtccgg 60  
 anttcccggg tcgaccacg cgtccgccc atctcccaga tccaattcgc gagttctttn 120

tcctctgccg ccatggcgct ctctgtggag aagacctcgt ctggacggga gtacaaggtc 180  
aaggatctct cgcaggcgga cttcgggccgc ctcgagattg agctggccga ggtcgaaatg 240  
cccggcctca tggcgtgccg cgcgagttc ggcccgctca agcccttcgc cggcgctagg 300  
atctcggggt ctctccacat gaccatccag accgcgctc tcatcgagac cctcaccgcg 360  
ctcgggcgcc aggtccgctg gtgctcctgc aacatcttct ccacgcagga ccacgccgcc 420  
gccgccatcg cgcgcgactc ggccgccgtg ttgccttga agggggagac cctcgaggag 480  
tactggtggt gcacccgagc gctgnctnga ctggggcnag gcggggccg 528

<210> 1365  
<211> 354  
<212> nucleic acid  
<213> Zea mays

<400> 1365  
cgtgccgcgc cgagttcggc ccgtccaagc ctttcgccgg cgtaggatc tcggggtctc 60  
tccacatgac catccagacc gccgtcctca tcgagaccct caccgcgctc ggcgccgagg 120  
tccgctggtg ctcttgcaac atcttctcca cgcaggacca cgccgccgcc gccatcgcg 180  
gcgactcggc cgccgtgttc gcctggaagg gggagaccct cgaggagtac tgggtggtgca 240  
ccgagcgctg cctcgactgg ggcgaggcgg gcggccccga cctcatcgtc gacgacggcg 300  
gcgacgccac gctgctcatc cacgagggtg tcaaggccga ggaggattac nana 354

<210> 1366  
<211> 476  
<212> nucleic acid  
<213> Zea mays

<400> 1366  
gggggngnaa cttcanntcc cagnngccgg tcaagaatta acgggtcgac gcncgcgtcc 60  
ggctcaaggc tgaccccaag aagtaccgca agatgaagga gaggcttgtc ggcgctctctg 120  
aggagaccac cacgggtgtc aagaggctct accagatgca ggagaccggc gccctcctct 180  
tccctgccat taacgtcaac gattccgtca ccaagagcaa gtttgacaac ctgtatggtt 240  
gccgccactc gtcctctgat ggtctgatga gggccactga cgttatgatc gccggaaagg 300  
ttgccgtggt ctgcggatac ggtgatgtcg gcaagggttg tgctgctgcc ctcaagcagg 360

ctggtgcccc tgtcattgtg accgagatcg accccatctg tgcctccag gctctgaatg 420  
 ganggtcttc aagtccttcc cttggaggac gttgtctcct gaagctgnca tcttng 476

<210> 1367  
 <211> 468  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1367

atcatggttg accacatgag gaagatgaag aacaatgcc ttgtctgcaa cattggccac 60  
 tttgacaatg aaattgatat gctcggcctt gagacctacc ctggcgtcaa gcgcatcacc 120  
 atcaagcccc agactgaccg ctgggtgttc cccgagacca aactggcat cattgtcctt 180  
 gctgaggggtc gctgatgaa ccttgggtgt gctactggcc atcctagctt tgtcatgtcc 240  
 tgctcattca ctaaccaggt cattgcccaa cttgaactgt ggaaggagaa gagctctggc 300  
 aagtatgaga agaaggtgta tgtgcttccc aagcaccttg atgagaaagg tgctgtcttc 360  
 acttggggca agctttgggtg ccaagcttac caaacttaac caagtcttaa ggccgactac 420  
 attaaccgtg ccgcatcgaa gggtccttac aagcctgcc actnaccg 468

<210> 1368  
 <211> 484  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1368

tttctttgcc ctggccggta ccanatcana gaattcccgg gncgaccac gcattcngct 60  
 caaggctgac cccaagaagt accgcaagat gaaggagagg cttgtcggcg tctctgagga 120  
 gaccaccacg ggtgtcaaga ggctctacca gatgcaggag accggcgccc tcctcttccc 180  
 tgccattaac gtcaacgatt ccgtcaccaa gagcaagttt gacaacctgt atggttgccg 240  
 cactcgctc cctgatggtc tgatgagggc cactgacgtt atgatcgccg gaaaggttgc 300  
 cgtgggtctgc ggatacgggtg atgtcggcaa gggttgtgct gctgccctca agcaggctgg 360  
 tgcccgtgtc attgtgaccg agatcgaccc catctgtgcc tccanggctc tgaatggaag 420  
 gtcttcangt ccttcccttg gaaggacgtt ggtctccgaa agctgacatc ttcgtgacna 480  
 acaa 484

<210> 1369  
 <211> 416  
 <212> nucleic acid  
 <213> Zea mays

<400> 1369

tgctcggcct tgagacctac cctggcggtca agcgcatcac catcaagccc cagactgacc 60  
 gctgggtggtt ccccgagacc aacactggca tcattgtcct tgctgaggggt cgctgatga 120  
 accttgggtg tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg 180  
 tcattgccc aactgaactg tggaaggaga agagctctgg caagtatgag aagaagggtgt 240  
 atgtgctccc caagcacctt gatgagaagg ttgctgctct ccacttgggc aagcttggtg 300  
 ccaagctgac caagctcacc aagtctcagg ccgactacat cagcgtgccg atcgaggggc 360  
 cctacaagcc tgccactacc ggtactaggc aagccaggca cacggcttgc agctna 416

<210> 1370  
 <211> 452  
 <212> nucleic acid  
 <213> Zea mays

<400> 1370

attgatatgc tcggccttga gacctaccct ggcgtaagc gcatcaccat caagccccag 60  
 actgaccgct ggggtgttccc cgagaccaac actggcatca ttgtccttgc tgaggggtgc 120  
 ctgatgaacc ttgggtgtgc tactggccat cctagctttg tcatgtcctg ctcatcact 180  
 aaccagggtca ttgcccact tgaactgtgg aaggagaaga gctctggcaa gtatgagaag 240  
 aagggtgatg tgctcccca gacacttgat gagaagggtg ctgctctcca cttggggcaa 300  
 gcttggtgcc aagctgacca agctcaccaa gtcttaagcc gactacatca gcgtgccgat 360  
 cganggtccc tacaagcctg ccactaccgg tactangcaa gccagcacac ggcttgagc 420  
 ttactcggcc cgttgtgtgc tatgaagttc ct 452

<210> 1371  
 <211> 481  
 <212> nucleic acid  
 <213> Zea mays

<400> 1371



gcaagatgaa ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggn 60  
 nctaccagat gcaggagacc ggcgccctcc tnttcctgc cattaacgtc aacgattccg 120  
 tcaccaagag caagtttgac aacctgtatg gntgccgcca ctgcctccct gatggtctga 180  
 tgagggccac tgacgttatg atcgccggtg aggntgctgn ggtctgcgga tacggngatg 240  
 tcggcaaggg ttgngctgct gccctcaagc aggctggtgc ccgtgtcatt gggaccgaga 300  
 tcgaccccat ctgngccctc caaggctctg atggagggnc ttcaggncct tcccttgagg 360  
 gacgnngctn tgaaactgac atcttcggga ccaccactgg caacaaggat atcatcangg 420  
 gtgaccacat gaaggaagat gaagaacaaa gccaanngcn gggaacaatn ggccaanttg 480  
 a 481

1372  
 445  
 nucleic acid  
 Zea mays  
 1372

<210> 1372  
 <211> 445  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1372  
 cgccgtcttc atcgagaccc tcaccgcgnt cggcgccgag gtccgctggt gtccttcaa 60  
 catcttctcc acgcaggacc acgccgccgc cgccatcgcg cgcgactcgg ccgcctgtt 120  
 cgcttgaag ggggagaccc ttgaggagta ctggtggtgc accgagcgtt gcctcgactg 180  
 gggcgaggcg ggcggccccc acctcatcgt cgacgacggc ggcgacgcca cgctgctcat 240  
 ccacgagggt gtcaaggccg aggaggacta cgagaagacc ggcaagatcc ccgatccgga 300  
 gtcaccgaca acgctgagtt caagatcgtg ctnaccatca ttccgcgaan ggctnaaagn 360  
 ttaccccaag aagtcccga aaggataaag aaaagcttgt cggcgtnttt taagagacca 420  
 cacnggtgtn aaaaggnttt acaga 445

<210> 1373  
 <211> 512  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1373

ggggttggtg nnnnggtggg naaattatnt gttgtaccgg tccggaattc ccgggtcgac 60  
 ccacgcgtcc gccacgcgt ccgtcacgc gttcgccan nntccagat ncaattcgcg 120

agttctgggn tgctctgccg ccatggcgct ctctgtggag aagacctcgt ctggacggga 180  
gtacaaggtc aaggatctct cgcaggcgga ctccggctgc ctcgagattg agctggccga 240  
ggtcgaaatg cccggcctca tggcgctgcc cgccgagttc ggcccgtcca agcccttcgc 300  
cggcgctagg atctcggggg ctcttcacat gaccatccag accgccgtcc tcatcgagac 360  
cctcaccgcg ctcggcgcgc aggtccgctg gtgctcctgc aacatcttct ncacgcagga 420  
ccacgccgtc ggcggcacgc ngcgcgactc gngcggcggt ttcgcctgga agggggagac 480  
cctttganga gtactggtgg tncaccnaac cg 512

<210> 1374  
<211> 523  
<212> nucleic acid  
<213> Zea mays

<400> 1374

agaggctngt tttgantggg gattaanggn ncttaatgg aaagctcttc cggnccggn 60  
nccccgctcg acccangenn tncanntnt ggaggggtctt naggtcctnc ncttgaggga 120  
cgttgtctnn gnagctgaen gnttcgtgac caccactggc aacaaggata tcatcatggt 180  
tgaccacatg aggaagatga agaacaatgc cattgtctgc aacattggcc actttgacaa 240  
tgaaattgat atgctcggcc ttgagacctt ccctggcgct aagcgcatna ccatcaagcc 300  
ccagactgac cgctgggtgt tccccgagac caacactggc atcattgtcc ttgctgaggg 360  
tcgcctgatg aaccttgggt gtgctactgg ccctcctagc tttgtcatgt cctgctcatt 420  
cactaaccag gtcattgccc aacttgaact gtggaaggag aagagctctg gcaagtatga 480  
gaaagaangt gtatgtgctt cccaacacct ttgatgagaa ggt 523

<210> 1375  
<211> 441  
<212> nucleic acid  
<213> Zea mays

<400> 1375

cccagatcca attcgcgagt tctccctcct ctgccgccat ggcgctctct gtggagaaga 60  
cctcgtctgg acnggagtac aaggtaagg atctctcgca ggcggacttc ggccgcctcg 120  
agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gaggctcgcc 180

cgccaagcc cttcgccggc gctaggatct cgggggtctct ccacatgacc atccagaccg 240  
 ccgtctcat cgagaccctc accgcgctcg gcncccgagg tccgctgggtg ctcttgcaac 300  
 atctttctnca cgcaggacca cnccttcgnc ngcatcgcg cgcactcgnc cgcctgtgttc 360  
 gcctggaaag ggggagnccc tcnaggagta ctgggtgttc accnagccgc ttgccttgnc 420  
 tggggccnag gcngnccgnc c 441

<210> 1376  
 <211> 488  
 <212> nucleic acid  
 <213> Zea mays

<400> 1376

ggnnnnnngg naacttnaac tncagcgcc aggtaacggt caaagaattc ccgggtcgac 60  
 cacgcgtccg cggacgcgtg ggcccgcttc atttcccat ctcccagatc caattcgag 120  
 ttctccctcc tctgcccga tggcgctctc tgtggagaag acctcgctctg gacgggatac 180  
 aaggtcaagg atctctcgca ggcggacttc ggccgcctcg agattgagct ggccgagtcg 240  
 aaatgcccgg cctcatggcg tgccgcgcg agttcgggcc gtccaagccc ttcgcgcgcc 300  
 taggatctcg gggctctctc acatgaccat ccagaccgcc gtcctcatcg agacctcacc 360  
 gcgctcgggc cggaggtccg ctgggtgctc tgcaacatct tctccacgca aggccacgcc 420  
 gccgcgcga tcgcgcgcga ctggngcgc gtgttcgcct ggaaggggga gactcgaagg 480  
 attatggg 488

<210> 1377  
 <211> 325  
 <212> nucleic acid  
 <213> Zea mays

<400> 1377

caagaagtac cgcaagatga aggagaggct tgtcggcgct tctgaggaga ccaccacggg 60  
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 caacgattcc gtcaccaaga gcaagtttga caacctgtat ggttgccgcc actcactccc 180  
 tgatggtctg atgagggcca ccgacgttat gatcgccggt aaggttgccg tggctctgcg 240  
 atacggtgat gttggcaagg gttgtgccgc tgcactcaag caggctgggt cccgtgtcat 300

tgtgaccgag atcgacccca tctgc 325

<210> 1378  
 <211> 325  
 <212> nucleic acid  
 <213> Zea mays

<400> 1378

gaagctgaca ttttctgtgac caccactggc aacaaggata tcatcatggt tgaccacatg 60  
 aggaagatga agaacaatgc cattgtctgc aacattggcc actttgacaa tgaaattgat 120  
 atgctcggcc ttgagaccta ccctggcgtc aagcgcatca ccatcaagcc ccagactgac 180  
 cgtcgggtgt tccccgagac caacactggc atcattgtcc ttgctgaggg tcgcctgatg 240  
 aacottgggt gtgctactgg ccctcctagc tttgtcatgt cctgtcatt cactaaccag 300  
 gtcattgccc aacttgaact gtgga 325

<210> 1379  
 <211> 370  
 <212> nucleic acid  
 <213> Zea mays

<400> 1379

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 gagaagaccg gcaagatccc cgaccggag tccaccgaca acgctgagtt caagatcgtg 180  
 ctcaccatca tccgcgacgg gctcaaggct gacccaaga agtntcga gatgaaggag 240  
 aggtttgtcg gcgtctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag 300  
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 tgacaacctg 370

<210> 1380  
 <211> 329  
 <212> nucleic acid  
 <213> Zea mays

<400> 1380

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ctgacgttat gatcgccgga aagggtgccc tggtctgcgg atacggtgat gtcggcaagg 120  
gttgtgctgc tgccctcaag caggctgggtg cccgtgtcat tgtgaccgag atcgacccca 180  
totgtgccct ccaggctctg atggaggggc ttcaggtcct tcccttgag gacgttgtct 240  
ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg gttgaccaca 300  
tgangaagat gaagaacaat gccattgtc 329

<210> 1381  
<211> 346  
<212> nucleic acid  
<213> Zea mays

<400> 1381

cacgggtgtc aagaggctct accagatgca ggagaccggc gccctcctct tccctgccat 60  
taacgtcaac gattccgtca ccaagagcaa gtttgacaac ctgtatgggt gccgccactc 120  
actccctgat ggtctgatga gggccaccga cgttatgatc gccggtaagg ttgccgtgggt 180  
ctgoggatac ggtgatgttg gcaaggggtg tgccgctgca ctcaagcagg ctggtgcccg 240  
tgtcatgtga ccgagatcga ccccatctgc gccctccagg ctctgatgga ggggtcttcag 300  
tccttccctt ggaggacgtt gtctcggaag ctgacatctt cgtgac 346

<210> 1382  
<211> 320  
<212> nucleic acid  
<213> Zea mays

<400> 1382

ccgagcgtg ccttgactgg ggcgagggcg gcggccccga cctcatcgtc gacgacggcg 60  
ggaagccac gctgctcatc cacgaggggtg tcaaggccga ggaggagtac gagaagaccg 120  
gcaagatccc cgacnccgag tccaccgaca acgctgagtt caagatcgtg ctcaccatca 180  
tccgcgacgg gctcaaggct gaccccaaga agtaccgcaa gatgaaggag aggcttgtcg 240  
ggtctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag gagaccggcg 300  
ccctcctctt cctgccatt 320

<210> 1383  
<211> 455  
<212> nucleic acid

<213> Zea mays

<400> 1383

ctcaaggctg accccaagaa gtaccgcaag atgaaggaga ggcttgtcgg cgtctctgag 60  
gagaccacca cgggtgtcaa gaggtcttac cagatgcagg agaccggcgc cctcctcttc 120  
cctgccatta acgtcaacga ttccgtcacc aagagcaagt ttgacaacct gtatggttgc 180  
cggcactcgc tcctgatgg tctgatgagg gccactgacg ttatgatcgc cggtaagggt 240  
gctgtgggtct gcggatacng tgatgtcggc aaggggtgtg ctgctgcctn aaacaggctg 300  
gtgccccgtg tcattgtgac ccagatcgac cccatctgtg cccttcaagc ttctgatnga 360  
nngncttcan gtccttcctt tggaaggacg ttgtntttgn aacttgacat ttttngntgg 420  
accaccactt gggaacaagg ggtnttatta ttggg 455

<210> 1384

<211> 317

<212> nucleic acid

<213> Zea mays

<400> 1384

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ggaggacgtt gtctctgaag ctgacatctt cgtgaccacc actggcaaca aggatatcat 120  
catggttgac cacatgagga agatgaagaa caatgccatt gtctgcaaca ttggccactt 180  
tgacaatgaa attgatatgc tcggccttga gacctaccct ggcgtcaagc gcatcaccat 240  
caagccccag actgaccgct ggggtgttccc cgagaccaac actggcatca ttgtccttgc 300  
tgagggtcgc ctgatga 317

<210> 1385

<211> 332

<212> nucleic acid

<213> Zea mays

<400> 1385

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cccagacca aactggcat cattgtcctt gctgagggtc gcctgatgaa cttgggtgt 180

getactggcc atcctagctt tgtcatgtcc tgctcattca ctaaccaggt cattgccc aa 240  
 cttgaactgt ggaaggagaa gagctctggc aagtatgaga agaagggtga tgtgctcccc 300  
 aagcaccttg atgagaaggt tgctgctctc ca 332

<210> 1386  
 <211> 337  
 <212> nucleic acid  
 <213> Zea mays

<400> 1386

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 gatgcgcggt aaggttgccg tgggtctgcg atacgggtgat gttggcaagg gttgtgcgcg 180  
 tgcaactcaag caggctgggtg cccgtgtcat tgtgaccgag atcgacccca tctgcgcctt 240  
 ccaggctctg atggaggggtc ttcaggctct tcccttgag gagttgtctc ggaagctgac 300  
 atcttcgtga ccacccatgg caacaaggat atcatca 337

<210> 1387  
 <211> 316  
 <212> nucleic acid  
 <213> Zea mays

<400> 1387

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 ctggcaacaa ggatatcatc atgggtgacc acatgaggaa gatgaagaac aatgccattg 120  
 tctgcaacat tggccacttt gacaatgaaa ttgatatgct cggccttgag acctaccctg 180  
 gcgtcaagcg catcaccatc aagccccaga ctgaccgctg ggtgttcccc gagaccaaca 240  
 ctggcatcat tgtccttgct ganggtcgcc tgatgaacct tgggtgtgct actggccatc 300  
 ctagctttgt catgtc 316

<210> 1388  
 <211> 315  
 <212> nucleic acid  
 <213> Zea mays

<400> 1388

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gcccgcgcca tncgcgcna ctcgcccgcc gtgttcgcct ggaaggggga gacccttgan 120  
gagtactggg ggtgcaccga gcgctgcctt gactggngcg angcggggcg ccccnacctc 180  
atcgctgcag acggcggcga cgccacgctg ctcatccaag aggggtgtcaa ggccgaggag 240  
gagtacgaga agaccggcaa gatccccgac ccggagtcca ccgacaacgc tgagttcaag 300  
atcggtgtca ccatac 315

<210> 1389  
<211> 310  
<212> nucleic acid  
<213> Zea mays

<400> 1389

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ggttgcgcgc actcgctccc tgatggtctg atgagggcca ctgacgttat gatcgccgga 180  
aaggttgcgc tgggtctgcg atacggtgat gtcggcaagg gttgtgctgc tgccctcaag 240  
caggctggtg cccgtgtcat tgtgaccgag atcgacccca tctgtgccct ccaggctctg 300  
atggaggggc 310

<210> 1390  
<211> 457  
<212> nucleic acid  
<213> Zea mays

<400> 1390

gggtctctc cacatgacca tccagacnng caagatcccc gatccggagt ccaccngna 60  
cgctgagttc aagatcgtgc tcaccatcat ccgcgacggg ctcaaggctg accccaagan 120  
gtaccgcaag atgaaggaga ggcttgctcg cgtctntgag gagaccacca cgggtgtcaa 180  
gaggctctac cagatgcagg agaccggcgc cctcctcttc ctgccattaa cgtcaacgat 240  
tccgtcacca agagcaagtt tgacaacctg tatggttgcc gncactcgct ccctgatggt 300  
ctgatgaagg gccactgacc ttatgatcgc ccgaaanggt gccgtggtct gcggataccg 360  
tgatgtcngc aaagggttgt gcttnttnan ttaaancang cttggtggcc ctgtcantnt 420



gaaccananc caancccatn ttggncctt cagggtt

457

<210> 1391  
<211> 520  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1391

agtttattna aggnngngg ntgnaagnta tactcgtncg gaattcccg gtcgacccac 60  
gogtccgata tencagaccc aattcgngag ttctnnntan tctgccgaca tggcgctctc 120  
tgtggagaag acctgntctg gacgggagta caaggtcaag gatctctagc aggcggactn 180  
aggccgacta gagattgagc tggccgaggt cgaaatgcc ggccatnatgg cgtgccgngc 240  
cgagttcggc ccgtncgaagc ccttngccgg cgctaggatc tcgggggtctc tccacatgac 300  
cattcagacc gncgtcctca tcgagaccct caccgcgctc ggcgccgagg tccgctggtg 360  
ctcctgcaac atctttcttc acgcangacc acgccgncgn cgtcatcgcn cgcgactcgg 420  
ccggcggtgtt cgcttggaaan ggggagaccc ttgangagtc tgggtggtgca ccgnacgctt 480  
gcntanntgg gccaaagcggc cggcccgacc tattgtngac 520

<210> 1392  
<211> 305  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1392

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ttacgagaag accggcaaga tctccgaccc ggagtccacc gacaacgctg agttcaagat 120  
cgtgctcacc atcatccgag acgggctcaa ggctgacccc aagaagtacc gcaagatgaa 180  
ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 240  
gcaggagacc ggccgccctcc tcttccctgc cattaacgct aacgattccg tcaccaagag 300  
caagt 305

<210> 1393  
<211> 317  
<212> nucleic acid  
<213> Zea mays

<400> 1393

gtgcccgtgt cattgtgacc gagatcgacc ccatctgcnc cctccaggct ctgatggagg 60  
gttttcagggt ctttcccttg gaggacgttg tctcggaagc tgacatcttc gtgaccacca 120  
ctggcaacaa ggatatcatc atggttgacc acatgaggaa gatgaagaac aatgccattg 180  
totgcaacat tggccacttt ganaatgaaa ttgatatgct cggccttgag acctaccctg 240  
gcgtaagcg catcaccatc aagccccaga ctgaccgctg ggtgttcccc gagaccaaca 300  
ctggcatcat tgtcctt 317

<210> 1394

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 1394

gctgccttga ctggggcgag gcgggcggcc ccgacctcat cgtcgacgac ggcggcgacg 60  
ccacgctgct catccacgag ggtgncaagg ccgaggagga gtacgagaag accggcaaga 120  
tccccgaccc ggagtccacc gacaacgctg agttcaagat cgtgctcacc atcatccgcg 180  
acgggctcaa ggctgacccc aagaagtacc gcaagatgaa ggagaggctt gtcggcgtct 240  
ctgaggagac caccacgggt gtcaagaggc tctaccagat gcaggagacc ggcgccctcc 300  
tcttccctg 309

<210> 1395

<211> 335

<212> nucleic acid

<213> Zea mays

<400> 1395

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gctgagggtc gctgatgaa cttgggtgt gctactggcc atcctagctt tgtcatgtcc 120  
tgtcattca ctaaccagggt cattgcccaa cttgaactgt ggaaggagaa gagctctggc 180  
aagtatgaga agaaggtgta tgtgctcccc aagcaccttg atgagaaggt tgtgctctc 240  
caattgggca agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc 300  
agcgtgccga tcgaggggtcc ctacaagcct gccca 335



caacctgtat ggttgccgcc actcgctccc tgatggtctg atgagggcca ctgacgttat 240  
 gatcgccgga aaggttgccg tggctctgcgg atacggtgat gtcggcaagg gttgtgctgc 300  
 tgc 303

<210> 1399  
 <211> 311  
 <212> nucleic acid  
 <213> Zea mays

<400> 1399

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 tgaccaccac tggcaacaag gatatcatca tggttgacca catgaggaag atgaagaaca 180  
 atgccattgt ctgcaacatt ggccactttg acaatgaaat tgatatnctc ggcttgaga 240  
 cctaccctgg cgtcaagcgc atcaccatca agccccagac tgaccgctgg gtgttccccg 300  
 agaccaacac t 311

<210> 1400  
 <211> 308  
 <212> nucleic acid  
 <213> Zea mays

<400> 1400

caaggatata atcatggttg accacatgag gaagatgaag aacaatgcca ttgtctgcaa 60  
 cattggccac tttgacaatg aaattgatat gtcggcctt gagacctacc ctggcgtcaa 120  
 ggcatacacc atcaagcccc agactgaccg ctggngttc cccgagacca aactggcat 180  
 cattgtcctt gctgagggtc gctgatgaa cttgggtgt gctactggcc atcctagctt 240  
 tgtcatgtcc tgctcattca ctaaccaggt cattgccc aa cttgaactgt ggaaggagaa 300  
 gagctctg 308

<210> 1401  
 <211> 309  
 <212> nucleic acid  
 <213> Zea mays

<400> 1401

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 ggagggtott cagggtccttc ccttgaggga cgttgtctcg gaagctgaca tcttcgtgac 120  
 caccactggc aacaaggata tcatcatggt tgaccacatg aggaagatga agaacaatgc 180  
 cattgtctgc aacattggcc actttgacaa tgaaattgat atgctcggcc ttgagacctg 240  
 ccctggcgtc aagcgcatca ccatcaagcc ccagactgac cgctgggtgt tccccgagac 300  
 caacactgg 309

<210> 1402  
 <211> 311  
 <212> nucleic acid  
 <213> Zea mays

<400> 1402

ctctgtgacc accactggca acaaggatat catcatgggt gaccacatga ggaagatgaa 60  
 gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcggcct 120  
 tgagacctac cctggcgta agcgcatcac catcaagccc cagactgacc gctgggtggt 180  
 ccccgagacc aacaactggca tcattgtcct tgctgagggt cgctgatga accttgggtg 240  
 tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg tcattgcca 300  
 cttgaactgt g 311

<210> 1403  
 <211> 338  
 <212> nucleic acid  
 <213> Zea mays

<400> 1403

gtttgacaac ctgtatggtt gccgccactc gctccctgat ggtctgatga gggccactga 60  
 cgttatgatc gccggaaang ttgccgtggt ctgcggatac ggtgatgtcg gcaagggttg 120  
 tgctgctgcc ctcaagcagg ctggtgcccg tgtcattgtg accgagatcg acccatctg 180  
 tgccctccag gctctgatgg agggctttca ggtccttccc ttggaggacg ttgtctctga 240  
 agctgacatc ttctgaccca ccaactggcaa caaggatatc atcagggtga ccacatgang 300  
 aagatgaaga acaatgccat gtctgcaaca tggccant 338

<210> 1404

<211> 306  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1404  
  
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 ccccgagacc aacactggca tcattgtcct tgctgagggt cgctgatga accttgggtg 120  
 tgctactggc catcctagct ttgtcatgtc ctgtcattc actaaccagg tcattgccc 180  
 acttgaactg tggaaggaga agagctctgg caagtatgag aagaagggtg atgtgctccc 240  
 caagcacctt gatgagaagg ttgctgctct ccaattgggc aagcttggtg ccaagctgac 300  
 caagct 306

<210> 1405  
 <211> 424  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1405  
  
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 ancaanntga ngcccagatc attgtnaccg agatcgaccc catctgtgcc ctncaggctc 120  
 tgatggaggg tcttcaggct cttcccttgn aggacgttgt ctctgaagct gacatcttcg 180  
 tgaccaccac tggcaacaag gatatcatca tggttgacca catgaggaag atgaagaaca 240  
 atgccattgt ctgcaacatt ggccactttg acaatgaaat tgatatgctc ggccttgaga 300  
 cctaccctgn cgtcaagcgc atcaccatca agccccagac tgaccgctgg gtgttccccg 360  
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 gcta 424

<210> 1406  
 <211> 299  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1406  
  
 gtgaccgaga tcgaccccat ctgtgccctc caggctctga tggagggtct tcaggctcct 60  
 cccttgaggg acgttgtctc tgaagctgac atcttcgtga ccaccactgg caacaaggat 120

atcatcatgg ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc 180  
 cactttgaca atgaaattga tatgctcggc cttgagacct accctggcgt caagcgcac 240  
 accatcaagc cccagactga ccgctgggtg ttccccgaga ccaacactgg catcattgt 299

<210> 1407  
 <211> 299  
 <212> nucleic acid  
 <213> Zea mays

<400> 1407

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 ccgcgcctct catcgagacc ctcaccgcgc tcggcgccga ggtccgctgg tgctcctgca 120  
 acatcttctc cagcgaggac cagcgccgcg ccgccatcgc gcgcnactcg gccgcgctgt 180  
 tcgcctggna gggggagacc ctcgaggagt actggtggtg caccgagcgc tgctctgact 240  
 ggngcgangc gggcggnccc gacctcatcg tcgacgacgg cggcgacgcc acgtgtctc 299

<210> 1408  
 <211> 303  
 <212> nucleic acid  
 <213> Zea mays

<400> 1408

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 gggggagacc cttgaggagt actggtggtg caccgagcgc tgctctgact ggggcgagggc 120  
 gggcgggccc gacctcatcg tcgacgacgg cggcgacgcc acgtgtctca tccacgaggg 180  
 tgtcaaggcc gaggaggagt acgagaagac cggcaagatc cccgaccgg agtccaccga 240  
 caacgtgag ttcaagatcg tgctcaccat catccgcgac gggctcaagg ctgaccccaa 300  
 gaa 303

<210> 1409  
 <211> 494  
 <212> nucleic acid  
 <213> Zea mays

<400> 1409

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ccanttacaa ttccncatct cccatgattc aatttcgcga agttctccct cctctgcccc 120  
atggcgctct ctgtgganaa gacctcgtct ggacgggagt acaagggtcaa ggatctccgc 180  
angcggactt cggncgcctc gagattgagc tggccgaggt cgaaatgccc ggctcttggc 240  
gttgccgcgc cgagttcggc cgtcnaagc cttcgtctgg cgctaggatc tcgggtctct 300  
ccacatgacc atccaaaccg cgtcctcat cgagaccctc accgcgctcg gcgcgaggtc 360  
cgctggtgct cctgcaacat cttctccacg cagaccacg ccgccgccgc catgcgcgcg 420  
actcgccgcg cgntgttcnc cctggaangg gggaaaacct ccaagaanta ctgtggttca 480  
ancgagccgc tgnt 494

<210> 1410  
<211> 299  
<212> nucleic acid  
<213> Zea mays

<400> 1410  
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ccatcgcgcg cgantcggcc gcngtgtncg cctggaaggg ggagaccctc gangagtact 180  
ggnggtgcac cgagcgctgc ctgcactggn gcgangcggg cggccccgac ctcatcgctg 240  
acgacggcgg cgacgccacg ctgctcatcc acgagggtgt caangccgag gaggattac 299

<210> 1411  
<211> 302  
<212> nucleic acid  
<213> Zea mays

<400> 1411  
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cgtgctcacc atcatccgcg acggggtcaa ggctgacccc aagaagtacc gcaagatgaa 180  
ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 240  
gcagganacc ggccgacctc tcttcctgc cattaacgtc nacgattccg tcaccaagag 300  
ca 302



<210> 1412  
 <211> 485  
 <212> nucleic acid  
 <213> Zea mays

<400> 1412

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 ntntgtggag aagacotcgt ctggacggga gtacaaggtc aaggatctct cgcaggcgga 120  
 cttcggtcgc ctgagattg agctggccga ggtcgaaatg cccggcctca tggcgtgccg 180  
 cgccgagttc ggcccgtcca agcccttcgc cggcgctagg atctcgggggt ctcttcacat 240  
 gaccatccag accgncgtcc tcatcgagac cctcaccng ctcggcgccg aggtccgctg 300  
 gtgctctgca acatcttntt cagcgaagga cagccgncg gccgncatcg cgcgcgactc 360  
 ggccggcgng ttgcctgga aagggggaga ccctttgagg agtactggtg gtgcaccgag 420  
 ccgcttgnet tganttgagg ccaggcnggg cggcccgaac ctnaatggtn gacaacnggg 480  
 gggaa 485

<210> 1413  
 <211> 311  
 <212> nucleic acid  
 <213> Zea mays

<400> 1413

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 tgctcattca ctaaccaggt cattgcccga cttgaactgt ggaaggagaa gagctctggc 180  
 aagtatgaga agaaggtgta tgtgctcccc aagcaccttg atgagaaggt tgctgctctc 240  
 cacttgggca agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc 300  
 agcgtgccga t 311

<210> 1414  
 <211> 311  
 <212> nucleic acid  
 <213> Zea mays

<400> 1414

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ggttgccgcc actcgctccc tganggtctg atgagggcca ctgacgttat gatcgccgga 180  
aaggttgcng tggnttgcgg atacggtgat gtcggcaagg gttgtgctgc tgccctcaag 240  
caggctggtg cccgtgtcat tgtgaccgan atnacccca tctgtgctc caggctctga 300  
tggaggggtct t 311

<210> 1415  
<211> 280  
<212> nucleic acid  
<213> Zea mays

<400> 1415

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cgccgcgcgc atcgcgcgcg actcgggcgc cgtgttcgcc tggaaggggg agacctcga 120  
ggagtaactgg tgggtgcaccg agcgtgcct cgactggggc gaggcgggcg gccccgacct 180  
catcgctgac gacggcgggc acgccacgct gctcatccac gagggtgtca aggccgagga 240  
ggattacgag aagaccggca agatccccga cccggagtcc 280

<210> 1416  
<211> 295  
<212> nucleic acid  
<213> Zea mays

<400> 1416

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gaggacgttg tctcggaagc tgacatcttc gtgaccacca ctggcaacaa ggatatcatc 120  
atggttgacc acatgaggaa gatgaagaac aatgccattg tctgcaacat tggccacttt 180  
gacaatgaaa ttgatatgct cggccttgag acctaccctg gcgtcaagcg catcaccatc 240  
aagccccaga ctgaccgctg ggtgttcccc gagaccaaca ctggcatcat tgtcc 295

<210> 1417  
<211> 349  
<212> nucleic acid  
<213> Zea mays

<400> 1417

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aactggcat cattgtcctt gctgaggtc gcctgatgaa ccttgggtgt gctactggcc 120  
atcctagctt tgtcatgtcc tgetcattca ctaaccaggt cattgccccaa cttgaactgt 180  
ggaaggagaa gagctctggc aagtatgaga agaaggtgta tgtgtctccc aagcaccttg 240  
atgagaaggt tgctgctctc cacttgggca agcttgggtgc caagctggac caagctcacc 300  
aagtctcagg ccgatacatc agcgtgccga tcgnggtcct acaagcctg 349

<210> 1418

<211> 292

<212> nucleic acid

<213> Zea mays

<400> 1418

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gatgttggca agggtttgtgc cgctgcactc aagcaggctg gtgcccggtg catttgtgacc 120  
gagatcgacc ccattctgcgc cctccaggct ctgatggagg gtcttcagggt ccttcccttg 180  
gaggacgttg tctcggaagc tgacatcttc gtgaccacca ctggcaacaa ggatatcatc 240  
atggttgacc acatgaggaa gatgaagaac aatgccattg tctgcaacat tg 292

<210> 1419

<211> 287

<212> nucleic acid

<213> Zea mays

<400> 1419

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attccgtcac caagagcaag ttgacaacc tgtatgggtg ccgccactca ctccctgatg 120  
gtctgatgag ggccaccgac gttatgatcg ccgtaagggt tgccgtggtc tgcggatacg 180  
gtgatgttgg caagggttgt gccgtgcac tcaagcaggc tgggtgccgt gtcattgtga 240  
ccgagatcga ccccatctgc gccctccagg ctctgatgga gggctctt 287

<210> 1420

<211> 304

<212> nucleic acid

<213> Zea mays

<400> 1420

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aagctgacat cttcgtgacc accactggca acaangatat catcatggtt gaccacatga 120  
ggaagatgaa gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata 180  
tgctcggcct tgagacctac cctggcgctca agcgcatcac catcaagccc cagactgacc 240  
gctgggtggt ccccgagacc aacactggca tcatgtcttg ctganggtcg cctgatgaac 300  
cttg 304

<210> 1421

<211> 283

<212> nucleic acid

<213> Zea mays

<400> 1421

ggagtactgg tgggtgcaccg agcgctgcct cgactggggc gaggcgggcg gccccgacct 60  
catcgtcgac gacggcgggcg acgccacgct gtcctccac gaggggtgtca aggccgagga 120  
ggattacnag aagaccggca agatccccga cccggagtcc accgacaacg ctgagttcaa 180  
gatcgtgctc accatcatcc gcgacgggct caaggctgac cccaagaagt accgcaagat 240  
gaaggagagg cttgtcggcg tctctgagga gaccaccacg ggt 283

<210> 1422

<211> 420

<212> nucleic acid

<213> Zea mays

<400> 1422

ggatatgctc cgggcttgag acctaccctg gcgtaagcg catnaccatc aagccccaga 60  
ctgaccgctg ggtgttcccc gagaccaaca ctggcatcat tgncttctgct gagggctcgcc 120  
tgatgaacct tgggtgtgct actggccatn ctagctttgt catgtcctgc tcattcacta 180  
accaggtcat tgcccaactt gaactgtgga aggagaagag ctctggcaag tatnanaaga 240  
angtgtatgt gctncccaag caccttgatn agaangntgn tgntctncac ttgggcaagc 300  
ttggtgcaa nctnaccaag cttaccaaag tcttaagncc gctacnttaa cctgcccctc 360

gaaggntcct tccaacctnn ccacnaaccg gtcttagnaa gcnnnacaac ggttngaant 420

<210> 1423  
 <211> 311  
 <212> nucleic acid  
 <213> Zea mays

<400> 1423

gaccggcgcc ctctcttccc ctgccattaa cgtcaacgat tccntcacca agancaagtt 60

tgacaacctg tatggttnen gccactcgct cctgatggg ctgatgagg ccactgacgt 120

tatgatcgcc ggaaagggtg cgtgggtctg cngatacgg gatgtcggca aggnttggtg 180

tgctgccctc aagcaggctg gtgcccggtg cattgtgacc ganatcgacc ccctctnttc 240

cctccaggct ctgatggagg gtcttcagg ccttccttg gaagacgttg tctctgaagc 300

tgacatcttc g 311

<210> 1424  
 <211> 283  
 <212> nucleic acid  
 <213> Zea mays

<400> 1424

cgacgccacg ctgctcatcc acgagggtgt caaggccgag gaggagtacg agaagaccgg 60

caagatcccc gaccgggagt ccaccgacaa cgctgagttc aagatcgtgc tcaccatcat 120

ccgcgacggg ctcaaggctg accccaagaa gtaccgcaag atgaaggaga ggcttgctcg 180

cgtctctgag gagaccacca cgggtgtcaa gaggctctac cagatgcagg agaccggcgc 240

cctctcttc cctgccatta acgtcaacga ttccgtcacc aag 283

<210> 1425  
 <211> 369  
 <212> nucleic acid  
 <213> Zea mays

<400> 1425

aattcctca cnaaaagcaa gtttganaac ctgtatggtt gccgnannc actccctgat 60

ggnetgatna gggccaccga cgttatgat gccgtaagg ttgccgtggn ctgcngatac 120

cgtgatgttg gcaanggttg tgccnctgca ctcaagcagg ctgntgcccg tgtcattgtg 180

accgagatcg annccatctg cgccctccac gctctgatgg atgggtcttc aagtccttcc 240  
 cttggaggac gttgtctcgg gaagctgaca tcttcgtgac caccactggc aacaaggata 300  
 tcatcaatgg gttgancaca tgaaggaacg atgaaggaca atggcantgt ctgcaacatt 360  
 gggcaacnt 369

<210> 1426  
 <211> 278  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1426

gcaagatccc cgaccggag tccaccgaca acgctgagtt caagatcgtg ctcaccatca 60  
 tccgcgacgg gctcaaggct gaccccaaga agtaccgcaa gatgaaggag aggcttgtcg 120  
 gcgtctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag gagaccggcg 180  
 ccctcctctt ccttgccatt aacgtcaacg attccgtcac caagagcaag tttgacaacc 240  
 tgtatgggtg ccgccactcg ctcctgatg gtctgatg 278

<210> 1427  
 <211> 275  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1427

cttcgtgacc accactggca acaaggatat catcatggtt gaccacatga ggaagatgaa 60  
 gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata tgctcggcct 120  
 tgagacctac cctggcgtca agcgcatcac catcaagccc cagactgacc gctgggtggt 180  
 ccccgagacc aacactggca tcattgtcct tgctgagggt cgcctgatga accttgggtg 240  
 tgctactggc catcctagct ttgtcatgtc ctgct 275

<210> 1428  
 <211> 275  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1428

tgacatcttc gtgaccacca ctggcaacaa ggatatcatc atggttgacc acatgaggaa 60

gatgaagaac aatgccattg tctgcaacat tggccacttt gacaatgaaa ttgatatgct 120  
 cggccttgag acctaccctg gcgtaagcg catcaccatc aagccccaga ctgaccgctg 180  
 ggtgttcccc gagaccaaca ctggcatcat tgtccttgct gagggtcgcc tgatgaacct 240  
 tgggtgtgct actggccatc ctagctttgt catgt 275

<210> 1429  
 <211> 294  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1429

caccatcaag cccagactg accgctgggt gttccccgag accaactctg gcatcattgt 60  
 ccttgctgag ggctgcctga tgaaccttgg gtgtgctact ggccatccta gctttgtcat 120  
 gtctgtctca ttactaacc aggtcattgc ccaacttgaa ctgtggaagg agaagagctc 180  
 tggcaagtat gagaagaagg tgtatgtgct cccaagcac cttgatgaga aggttgctgc 240  
 tctccacttg ggcaagcttg gtgccaagct gaccaagctc accaagtctc aggc 294

<210> 1430  
 <211> 276  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1430

gctgacatct tcgtgaccac cactggcaac aaggatatca tcatggttga ccacatgagg 60  
 aagatgaaga acaatgccat tgtctgcaac attggccact ttgacaatga aattgatatg 120  
 ctcggccttg agacctacc ttggcgtcaag cgcatacca tcaagcccca gactgaccgc 180  
 tgggtgttcc ccgagaccaa cactggcatc attgtccttg ctgagggtcg cctgatgaac 240  
 cttgggtgtg ctactggcca tcctagcttt gtcattg 276

<210> 1431  
 <211> 288  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1431

ctcatggcgt gccgcgccga gttcgggccc tccaagccct tcgccggcgc taggatctcg 60

gggtctctcc acatgaccat ccagaccgcc gtctctcatcg agaccctcac cgcgctcggc 120  
gccgaggtcc gctggtgctc ctgcaacatc ttctccacgc aggaccacgc cgccgccgcc 180  
atcgcgcgcg actcgccgc cgtgttcgcc tggnaagggg agacccttga ggagtactgg 240  
tggtgcaccg agcgctgcct tgactggggc gangcggggc gccccgac 288

<210> 1432  
<211> 285  
<212> nucleic acid  
<213> Zea mays  
<400> 1432

tgcaggagac cggcgccctc ctcttccctg ccattaacgt caacgattcc gtcaccaaga 60  
gcaagtttga caacctgtat ggttgccgcc actcgctccc tgatggtctg atgagggcca 120  
ctgacgttat gatcgccgga aaggttgccg tggctctgcg atacggtgat gtcggcaagg 180  
gttgtgctgc tgccctcaag caggctggtg cccgtgtcat tgtgaccgag atcgacccca 240  
tctgtgccct ccaggctctg atggaggggc ttcaggtcct tcct 285

<210> 1433  
<211> 280  
<212> nucleic acid  
<213> Zea mays  
<400> 1433

atcgacccca tctgtgccct ccaggctctg atggaggggc ttcaggtcct tcccttgag 60  
gacgttgtct ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg 120  
gttgaccaca tgaggaagat gaagaacaat gccattgtct gcaacattgg ccactttgac 180  
aatgaaattg atatgctcgg ccttgagacc taccctggcg tcaagcgcac caccatcaag 240  
ccccagactg accgctgggt gttccccgag accaacactg 280

<210> 1434  
<211> 316  
<212> nucleic acid  
<213> Zea mays  
<400> 1434

cgaccccatc tgcncctcc aggtctctgat ggagggctct caggtccttc ccttgaggga 60



cgttgtctcg gaagctgaca ttttcgtgac caccactggc aacaaggata tcatcatggt 120  
 tgaccacatg aggaagatga agaacaatgc cattgtctgc aacattggcc actttgacaa 180  
 tgaaattgat atgctcggcc ttgagacctc ccttggcgtc aagcgcatca ccatcaagcc 240  
 ccagactgac cgctgggtgt tccccgagac caacactggc atcatgtcct tgctgaaggt 300  
 cgctgatga acttgg 316

<210> 1435  
 <211> 298  
 <212> nucleic acid  
 <213> Zea mays

<400> 1435

gcccggcctc atggcggtgcc ggcgcgagtt cggcccgctc aagcccntcg cggcgctag 60  
 gatctcgggg tototccaca tgaccatcca gaccgccgtc ctcctcgaga cctcaccgc 120  
 gctcggcgcc gaggtccgct ggtgtccttg caacatcttc tccacgcagg accacgccgc 180  
 cgccgccatc ggcgcgcact cggccgcccgt gttcgcttgg aaagggggag accctcgagg 240  
 agtactggtg gtgcaccgag cgctgctcga ctggggcgaa gcggggcgcc cgacctca 298

<210> 1436  
 <211> 299  
 <212> nucleic acid  
 <213> Zea mays

<400> 1436

agaagaccgg caagatcccc gaccgggagt ccaccgacaa cgctgagttc aagatcgtgc 60  
 tcaccatcat ccgcgacggg ctcaaggctg accccaagaa gtaccgcaag atgaaggaga 120  
 ggcttgctcg cgtctctgag gagaccacca cgggtgtcaa gaggtcttac cagatgcagg 180  
 agaccggcgc cctcctcttc cctgccatta acgtcaacga ttccgtcacc aagagcaagt 240  
 ttgacaactg tatggttgcc gcanttcgtt ccttgatggt ttgatgaggg ccaactgang 299

<210> 1437  
 <211> 279  
 <212> nucleic acid  
 <213> Zea mays

<400> 1437

gtcaaggccg aggaggagta cgagaagacc ggcaagatcc ccgacccgga gtccaccgac 60  
aacgctgagt tcaagatcgt gctcaccatc atccgcgacg ggctcaaggc tgaccccaag 120  
aagtaccgca agatgaagga gaggtttgtc gggcgctctc gaggagacca ccacgggtgt 180  
caagaggctc taccagatgc aggagaccgg cgcctctctc ttccttgcca ttaacgtcaa 240  
cgattccgtc accaagagca agtttgacaa cctgtatgg 279

<210> 1438  
<211> 277  
<212> nucleic acid  
<213> Zea mays

<400> 1438

gcaagatccc cgacccggag tccaccgaca acgctgagtt caagatcgtg ctcaccatca 60  
tccgcgacgg gctcaaggct gaccccaaga agtaccgcaa gatgaaggag aggtttgtcg 120  
gcgctctctga ggagaccacc acgggtgtca agaggctcta ccagatgcag gagaccggcg 180  
ccctcctctt ccttgccatt aacgtcaacg attccgtcac caagagcaag ttgacaact 240  
gtatggttgc cgccaactgc tccctgatgg tctgatg 277

<210> 1439  
<211> 318  
<212> nucleic acid  
<213> Zea mays

<400> 1439

atttccccat ctcccagatc caattcgaga gttctccctc ctctgccgcc atggcgctct 60  
ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg caggcggact 120  
tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg gcgtgccgcg 180  
ccgagttcgg cccgtccaag cccttcgccg gcgctaggat ctcggggtct ctccacatga 240  
ccatccagac cgcctctctc atcgagaccc tcaccgcgtc cggcgccgag gtccgctggt 300  
gtcctgcaa catcttct 318

<210> 1440  
<211> 249  
<212> nucleic acid  
<213> Zea mays

<400> 1440

cacatgacca tccagaccgc cgtcctcatc gagaccctca ccgcgctcgg cgccgaggtc 60

cgctggtgct cctgcaacat cttctccacg caggaccacg ccgccgccgc catcgcgcg 120

gactcggccg ccgtgttcgc ctggaagggg gagaccctcg aggagtactg gtggtgcacc 180

gagcgtgcc tcgactgggg cgaggcgggc ggccccgacc tcatcgtcga cgacggcggc 240

gacgccacg 249

<210> 1441

<211> 309

<212> nucleic acid

<213> Zea mays

<400> 1441

cccagatcca attcgcgagt nonccctcct ctgcggccat ggcgctctct gtggagaaga 60

cctcgtctgg acgggagtac aagggtcaagg atctctcgca ggccggacttc ggccgcctcg 120

agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180

cgtccaagcc cttcgccggc gctaggatct cggggtctct ccacatgacc atccagaccg 240

ccgtcctcat cgagaccctn accgcgctcg gcgccgaggt ccgctggtgc tcctgcaaca 300

tcttctcca 309

<210> 1442

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1442

gtacgagaag accggcaaga tccccgaccc ggagtccacc gacaacgctg agttcaagat 60

cgtgctcacc atcatccgcg acggggtcaa ggctgacccc aagaagtacc gcaagatgaa 120

ggagaggctt gtcggcgtct ctgaggagac caccacgggt gtcaagaggc tctaccagat 180

gcaggagacc ggccgacctc tcttcctgc cattaacgtc aacgattccg tcaccaagag 240

caagtttgac aacctgtatg gttgccgcca ctact 276

<210> 1443

<211> 276

<212> nucleic acid

<213> Zea mays

<400> 1443

gtgaccgana tcgaccccat ctgtncctc caggctctga tggagggtct tcaggtcott 60  
cccttgaggg acgttgtctc tgaagctgac atnttcgtga ccaccactgg caacaaggat 120  
atcatcatgg ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc 180  
cactntgaca atgaaattga tatgctcggc cttgagacct accctggcgt caagcgcac 240  
accatcaagc cccagactga cgcgtgggtg ttcccc 276

<210> 1444

<211> 270

<212> nucleic acid

<213> Zea mays

<400> 1444

agaagaccgg caagatcccc gaccggagt ccaccgaca cgtgagttc aagatcgtgc 60  
tcaccatcat ccgcgacggg ctcaaggctg accccaagaa gtaccgcaag atgaaggaga 120  
ggcttgctgg cgtctctgag gagaccacca cgggtgtcaa gaggtcttac cagatgcagg 180  
agaccggcgc cctcctcttc cctgccatta acgtcaacga ttccgtcacc aagagcaagt 240  
ttgacaacct gtatggttgc cgccactcgc 270

<210> 1445

<211> 261

<212> nucleic acid

<213> Zea mays

<400> 1445

ggaggacgtt gtctctgaag ctgacatctt cgtgaccacc actggcaaca aggatatcat 60  
catggttgac cacatgagga agatgaagaa caatgccatt gtctgcaaca ttggccactt 120  
tgacaatgaa attgatatgc tcggccttga gacctacctt ggcgtaagc gcatcaccat 180  
caagccccag actgaccgct ggggtgtccc cgagaccaac actggcatca ttgtccttgc 240  
tgagggtcgc ctgatgaacc t 261

<210> 1446

<211> 291

<212> nucleic acid

<213> Zea mays  
 <400> 1446  
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 gttatgatcg ccggaaggt tgccgtggtc annccgatac ggtgatgtcg gcaagggttg 120  
 tgctgtgcc ctcaagcagg ctggtgcccg tgtcatttg accgagatcg accccatctg 180  
 tgccctccag gctctgatgg agggctctca ggtccttccc ttggaggacg ttgtctctga 240  
 agctgacatc ttcgtgacca ccactggcaa caaggatatc atcatggttg a 291

<210> 1447  
 <211> 316  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1447  
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 tcatccacga ggggtgtcaag gccgaggagg agtacgagaa gaccggcaag atccccgacc 120  
 cggagtccac cgacaacgct gagttcaaga tcgtgctcac catcatccgc gacgggctca 180  
 aggtgaccc caagaagtac cgcaagatga aggagagctt gtcggcgtct taaggagacc 240  
 accagggtgt caagaagctc taccagatgc aagaaaccgg cggcctcctc ttccctgcc 300  
 ttaacgtnac gatccg 316

<210> 1448  
 <211> 273  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1448  
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 gccgaggagg agtacgagaa gaccggcaag atccccgacc cggagtccac cgacaacgct 120  
 gagttcaaga tcgtgctcac catcatccgc gacgggctca aggtgaccc caagaagtac 180  
 cgcaagatga aggagaggct tgtcggcgctc tctgaggaga ccaccacggg tgncaagag 240  
 gctctaccag atgcaggaga cggcgccct cct 273

<210> 1449

<211> 271  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1449  
  
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 gcgcgcgaca cggccgccgt gttcgcttg aagggggaga ccctcgagga gtactggtgg 120  
 tgcaccgagc gctgcctcga ctggggcgag gcggggcgcc ccgacctcat cgtcgacgac 180  
 ggcggcgacg ccacgctgct catccacgag ggtgtcaagg ccgaggagga ttacgagaag 240  
 accggcaaga tccccgacct ggagtccacc g 271

<210> 1450  
 <211> 275  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1450  
  
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 gatggtctga tgagggccac tgacgttatg atcgccggaa aggttgccgt ggtctgcgga 120  
 tacggtgatg tcggcaaggg ttgtgctgct gccctcaagc aggttggtgc ccgtgtcatt 180  
 gtgaccgaga tcgaccccat ctgtgccttc caggctctga tggaggggtn caaggtcctn 240  
 cccttgaggg acgttgtctc ngaagatgac atctt 275

<210> 1451  
 <211> 271  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1451  
  
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 gaggacgttg tctcggaagc tgacatcttc gtgaccacca ctggcaacaa ggatatcatc 120  
 atggttgacc acatgaggaa gatgaagaac aatgccattg tctgcaacat tggccacttt 180  
 gacaatgaaa ttgatatgct cggccttgag acctaccctg gcgtcaagcg catcaccatc 240  
 aagccccaga ctgaccgctg ggtgttcccc g 271

<210> 1452



<211> 282  
 <212> nucleic acid  
 <213> Zea mays

<400> 1455

ccgccactcg ctccctgatg gtctgatgag ggccactgac gttatgatcg ccggaaaggt 60  
 tgccgtgggc tgcggatacg gtgatgtcnn gcaaggggtg tgctgctgcc ctcaagcagg 120  
 ctggtgcccc tgctattgtg accgagatcg accccatctg tgccctccag gctctgatgg 180  
 agggctttca ggctccttccc ttggaggacg ttgtctctga agctgacatc ttcgtgacca 240  
 ccaactggcaa caaggatata atcatgggtg accacatgag ga 282

<210> 1456  
 <211> 297  
 <212> nucleic acid  
 <213> Zea mays

<400> 1456

ggctgacccc aagaagtacc gcaagatgaa ggagaggctt gtcggcgtct ctgaggagac 60  
 caccacgggt gtcaagagggc tctaccagat gcaggagacc ggcgcctcc tcttccctgc 120  
 cattaacgtc aacgattccg tcaccaagag caagtttgac aacctgtatg gttgccgcca 180  
 ctgctccct gatggtctga tganggccac tgacgttatg attcgccgga aagggtgccg 240  
 ttgtctgcgg atacggtgat gtcggcaang gttgtgtgct gccctcaagc angctgg 297

<210> 1457  
 <211> 130  
 <212> nucleic acid  
 <213> Zea mays

<400> 1457

gngatcgacc ccatctgcgc cctccagnct ctnatggagg gtcttcaggt ccttcccttg 60  
 naggacgttg tctcggaagc tgacatcttc ggtgaccacc actggcaaca aggatatan 120  
 ncatgggttg 130

<210> 1458  
 <211> 304  
 <212> nucleic acid  
 <213> Zea mays



<400> 1458

catctcccag atccaattcg cgagttctcc ctctctgcg gccatggcgc tctctgtgga 60  
gaagacctcg tctggacggg agtacaaggt caaggatctc tcgcaggcgg acttcggccg 120  
cctcgagatt gagctggcgg aggtcgaaat gcccggcctc atggcgtgcc gcgccgagtt 180  
cgccccgtcc aagcccttcg cgggcgctag gatctcgggg tctctccaca tgaccatcca 240  
gaccgcgctc ctcatcgaga cctcaccgc gctcggcgcc gaggtccgct ggtgctcctg 300  
caac 304

<210> 1459

<211> 512

<212> nucleic acid

<213> Zea mays

<400> 1459

gnggnnnng agtgnntnnt aatttgagg naaaatgtaa nggnaatctt cgtaccggtc 60  
cggaannntc gaccacgcg tccgccacg cgtccggacc aacactggca tcattgtcct 120  
tgctgagggt cgnctgatga nctgggggtg tgctactggc catcctagct ttgncatgtc 180  
ctgtcattc actaaccagg tcattgcca acttgaactg tggaaggaga agagctctgg 240  
caagtatgaa aagaagggtg atgtntccc caagcacctt gatgagaagg ttgctgctct 300  
ccaattgggc aancttggtg ccaagctgac caagctnacc aagtctcagg ccgactacat 360  
cagcgtgccg atcgagggtc cctacaagcc tgccactacc ggtactaggc agccagcaca 420  
cggnttgcaa ctnactcggg cgtgtgtgc tatnaagccg ctactggcc tgnagntatc 480  
tnnognannc tatggcataa acatanacgg ga 512

<210> 1460

<211> 263

<212> nucleic acid

<213> Zea mays

<400> 1460

gccacgcagg accacgccgc cgccgccatc gcgcgcgact cggccgccgt gttcgctgn 60  
aanggggaga cccttgagga gtactgggtg tgnaccgagc gctgccttga ctggggcgag 120  
gcgggcggcc ccgacctcat cgtcgacgac ggccgacgac ccacgtgct catccaagag 180

ggtgtcaagg ccgaggagga gtacgagaag accggcaaga tccccgacct ggagtccacc 240  
gacaacgctg agttcaagat cgt 263

<210> 1461  
<211> 247  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1461

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ccgaggtcgg ctgggtgctcc tgcaacatct tctccacgca ggaccacgcc gccgcccga 120  
tcgcgcgcga ctgggcccgc gtgttcgcct ggaaggggga gacccttgag gagtactggt 180  
ggtgcaccga gcgctgcctt gactggggcg aggcggggcg ccccgacctc atcgtcgacg 240  
acggcgg 247

<210> 1462  
<211> 260  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1462

ggaagatgaa gaacaatgcc attgtctgca acattggcca ctttgacaat gaaattgata 60  
tgctcggcct tgagacctac cctggcgctca agcgcacac catcaagccc cagactgacc 120  
gtcgggtgtt ccccgagacc aacctggca tcattgtcct tgctgagggc cgctgatga 180  
accttgggtg tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg 240  
tcattgccc acttgaactg 260

<210> 1463  
<211> 272  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1463

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caagggttgt gccgctgcac tcaagcaggc tggcgcccg gtcattgtga ccgagatcga 120  
cccatctgc gccctccagg ctctgatgga gggctctcag gtccttcctc tggaggacgt 180

tgtctcggaa gctgacatct tcgtgaccac cactggcaac aaggatatca tcanggttga 240  
ccacatgagg aagatganga acaatgccat tg 272

<210> 1464  
<211> 253  
<212> nucleic acid  
<213> Zea mays

<400> 1464

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caatgaaatt gatatgctcg gccttgagac ctaccctggc gtcaagcgca tcaccatcaa 120  
gccccagact gaccgctggg tgttccccga gaccaacact ggcatcattg tccttgctga 180  
gggtcgctcg atgaaccttg ggtgtgctac tggccatcct agctttgtca tgtcctgctc 240  
attcactaac cag 253

<210> 1465  
<211> 261  
<212> nucleic acid  
<213> Zea mays

<400> 1465

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tccccgaccc ggagtccacc gacaacgctg agttcaagat cgtgctcacc atcatccgcg 120  
acgggctcaa ggctgacccc aagaagtacc gcaagatgaa ggagaggctt gtcggcgtct 180  
ctgaggagac caccacgggt gtcaagaggc tctaccagat gcaggagacc ggcgccctcc 240  
tcttccctgc cattaacgtc a 261

<210> 1466  
<211> 261  
<212> nucleic acid  
<213> Zea mays

<400> 1466

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ccatcgcgcg cgactcggcc gccgtgttcg cctggaaggg gagacccttg aggagtactg 180

gtggtgcacc gagcgtgcc ttgactgggg cgaggcgggc ggccccgacc tcatcgtcga 240  
cgacggcggc gacgcacgt g 261

<210> 1467  
<211> 323  
<212> nucleic acid  
<213> Zea mays  
<400> 1467

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atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaaggtaa ggatctctcg 120  
caggcggact tcggccgect cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180  
gcgtgccgcg ccgagttcgg ccggtccaag cccttcgccg ccgctaggat ctcggggtct 240  
ctccacatga ccatccagac cgcggtcctc atcgagacct tcaccgcgt cggcgccgag 300  
gtccgcaggt gtcctgcaa cat 323

<210> 1468  
<211> 277  
<212> nucleic acid  
<213> Zea mays  
<400> 1468

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gtcgggtgtt ccccgagacc aacctggca tcattgtcct tgctgagggt cgctgatga 120  
accttgggtg tgctactggc catcctagct ttgtcatgtc ctgctcattc actaaccagg 180  
tcattgcca acttgaactg tggaaggaga agagctctgg caagtatgag aagaagggtg 240  
atgtgctccc caagcacctt gatgagaagg ttgctgc 277

<210> 1469  
<211> 257  
<212> nucleic acid  
<213> Zea mays  
<400> 1469

caaggatatc atcatggttg accacatgag gaagatgaag aacaatgcca ttgtctgcaa 60  
cattggccac ttggaacaat gaaattgata tgctcggcct tgagacctac cctggcgtca 120

agcgcatcac catcaagccc cagactgacc gctgggtgtt ccccgagacc aacactggca 180  
tcattgtcct tgctgagggg cgctgatga accttgggtg tgctactggc catcctagct 240  
ttgtcatgtc ctgctca 257

<210> 1470  
<211> 262  
<212> nucleic acid  
<213> Zea mays

<400> 1470

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ggcgccctcc tcttccctgg ccattaacgt caacgattcc gtcaccaaga gcaagtttga 120  
caacctgtat gggtgccgcc actcgtccc tgatgggtctg atgagggcca ctgacgttat 180  
gatcgccgga aagggtgcg tggctctgcg atacggtgat gtcggcaagg gttgtgctgc 240  
tgcaactcaag caggctgggtg cc 262

<210> 1471  
<211> 317  
<212> nucleic acid  
<213> Zea mays

<400> 1471

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aagcaggctg gtgcccgtgt cattgtgacc gagatcgacc ccattctgnc cctccaggct 180  
ctgatggagg gtcttcagggt ccttcccttg gaggnntngt cacggaagct nanatttctg 240  
gaccaccact ggnaacaagg atatcatcat ggttgaccac atgaggaaga tgaanaacat 300  
gccattgtct cnaattg 317

<210> 1472  
<211> 268  
<212> nucleic acid  
<213> Zea mays

<400> 1472

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tgctcatcca cgaggggtgtc aaggccgagg aggattacga gaagaccggc aagatccccg 120  
 acccgaggtc caccgacaac gctgagttca agatcgtgct caccatcatc cgcgacgggc 180  
 tcaaggctga cccaagaag taccgcaaga tgaaggagag gcttgtcggc gtctctgagg 240  
 agaccaccac ggggtgtcaag aggtctcta 268

<210> 1473  
 <211> 274  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1473

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 gaccccatct gtgccctcca ggctctgatg gaggggtcttc aggtccttcc cttggaggac 120  
 gttgtctctg aagctgacat cttcgtgacc accactggca acaaggatat catcatgggt 180  
 gaccacatga ggaagatgaa gaacaatgcc attgtctgca acattggcca tttgacaatg 240  
 aaattgatat gctcggcctt gagacctacc ctgg 274

<210> 1474  
 <211> 290  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1474

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 tctgtgccct ccaggctctg atggaggggtc ttcaggctct tcccttgagg gacgttgtct 120  
 ctgaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg gttgaccaca 180  
 tgaggaagat gaagaacaat gccattgtct gcaacattgg ccactttgac aatgaaattg 240  
 atatgctcgg ccttgagact acctggcgtc aagcgcacat catcaagccc 290

<210> 1475  
 <211> 300  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1475

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caccgacaac gctgagttca agatcgtgct caccatcatc cgcgacgggc tcaaggctga 120  
 cccaagaag taccgcaaga tgaaggagag gcttgctggc gtctctgagg agaccaccac 180  
 ggggtgtcaag aggctctacc agatgcagga gaccggcgcc ctctcttcc ctgccattaa 240  
 cgtcaacgat tcgtcaccag agcaagtttg acnactgtat ggttgccgca attcattccc 300

<210> 1476  
 <211> 260  
 <212> nucleic acid  
 <213> Zea mays

<400> 1476

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 ccactggcaa caaggatata atcatgggtg accacatgag gaagatgaag aacaatgcc 120  
 ttgtctgcaa cattggccac ttgacaatg aaattgatat gtcggcctt gagacctacc 180  
 ctggcgtcaa ggcacatcac atcaagcccc agactgaccg ctgggtgttc cccgagacca 240  
 aactggcat cattgtcctt 260

<210> 1477  
 <211> 295  
 <212> nucleic acid  
 <213> Zea mays

<400> 1477

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 aggacgaggc ttgtcggcgt ctctgaggag accaccacgg gtgtcaagag gctctaccag 180  
 atgcaggaga ccggcgccct cctcttcct gccattaacg tcaangattc cgtcaccaag 240  
 agcaagtttg acaacntgta tggttgccgc caactcggct ccctgatggt ctgat 295

<210> 1478  
 <211> 278  
 <212> nucleic acid  
 <213> Zea mays

<400> 1478

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gaggagtact ggtggtgcac cgagcgctgc cttgactggg gcgaggcggg cggccccgac 120  
ctcatcgctcg acgacggcgg cgacgccaac gctgctcatc cacgagggtg tcaaggccga 180  
ggaggagtac gagaagaccg gcaagatccc cgacccggag tccaaccgac aacgntgagt 240  
tcaagatcgt gcttaccatc attcgggacn ggctcaaa 278

<210> 1479  
<211> 287  
<212> nucleic acid  
<213> Zea mays  
<400> 1479

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tgctcattca ctaaccaggt cattgcccac cttgaactgt ggaaggagaa gagctctggc 120  
aagtatgaga agaaggtgta tgtgctcccc aagcaccttg atgagaaggt tgctgctctc 180  
catttgggca agcttgggtgc caagctgacc aagctcacca agtctcaggc cgactacatc 240  
agcgtgccga tcgaggggtcc ctacaagcct gccactacc ggtacta 287

<210> 1480  
<211> 306  
<212> nucleic acid  
<213> Zea mays  
<400> 1480

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aatgagggcc actgacgtta tgatcgccgg aaaggttgcc gtggtctgcy gatacgggtga 120  
tntcggcaan ggttgtgctg ctgccctcaa gcaggctggt gcccggtgca ttgtgaccga 180  
gatcgacccc atctgtgccc tccaggctct gatggagggt cttcagggtcc ttcccttgga 240  
ggacgttgtc tctgaagctg acatcttcgt gaccaccact ggcaacaagg atatcacatg 300  
gttgac 306

<210> 1481  
<211> 314  
<212> nucleic acid  
<213> Zea mays  
<400> 1481



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catgaggaag atgaagaaca atgccattgt ctgcaacatt ggccactttg acaatgaaat 120  
tgatatgctc ggccttgaga cctaccctgg cgtcaagcgc atcaccatca agccccagac 180  
tgaccgctgg gtgttccccg agaccaacac tggcatcatt gtccttgctg anggtcgctt 240  
gatgaacttg ggtgtgtatg gccatcctag tttgtcatgt cctgtcatna ctaaccagtc 300  
attgnccaat tgaa 314

<210> 1482  
<211> 270  
<212> nucleic acid  
<213> Zea mays

<400> 1482  
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cctaccctgg cgtcaagcgc atcaccatac aagccccaga ctgaccgctg ggtgttcccc 120  
gagaccaaca ctggcatcat tgtccttgct gagggtcgcc tgatgaacct tgggtgtgct 180  
actggccatc ctagctttgt catgtcctgc tcattcacta accagggtcat tgcccaactt 240  
gaactgtgga aggagaagag ctctggcaag 270

<210> 1483  
<211> 266  
<212> nucleic acid  
<213> Zea mays

<400> 1483  
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ggnttggtgcc gctgcactca agcaggctgg tgcccgtntc attgtgaccg agatcgaccc 120  
catctgcnnn ctccangctc tgatggaggg tcttcaggtc cttcccttgg aggacgttgt 180  
ctoggaagct gacatcttcg tgaccaccac tggcaacaag gatatcatca tggttgacca 240  
catgaggaag atgaagaaca atgcca 266

<210> 1484  
<211> 312  
<212> nucleic acid  
<213> Zea mays

<400> 1484

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 ggcggaactt ggccgcctcg agattgagct ggccgaggtc gaaatgcccg gctcatggc 180  
 gtgccgcgnc gagttcggcc cgtccaagcc cttcgccggc gctaggatct cggggtctct 240  
 ccacatgacc atccagaccg cgtctctcat cgagaccctc accgcgctcg gcgccgaggt 300  
 ccgctggtgc tc 312

<210> 1485

<211> 271

<212> nucleic acid

<213> Zea mays

<400> 1485

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 tcattcacta accaggtcat tgcccaactt gaactgtgga aggagaagag ctctggcaag 180  
 tatgaaagaa ggtgtatgtg ctccccaagc accttgatga gaaggttgct gctctccact 240  
 tgggcaagct tggtgccaag ctgaccaagc t 271

<210> 1486

<211> 275

<212> nucleic acid

<213> Zea mays

<400> 1486

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 ccggagtcca ccgacaacgc tgagttcaag atcgtgtcga ccatcatccg cgacgggctc 180  
 aaggctgacc ccaagaagta ccgcaagatg aaggagaggc ttgtcggcgt ctctgaggag 240  
 accaccacgg gtgtcaagag gtctaccaga tgcag 275

<210> 1487

<211> 407

<212> nucleic acid

<213> Zea mays

<400> 1487

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ctctggcaag tatgagaaga aggtgtatgt gctccccaag caccttgatg agaaggttgc 180

tgctctccac ttgggcaagc ttggtgccaa gctgaccaag ctcaccaagt ctcaggccga 240

ctacatcagc gtgcgatcg agggtccta caagcctgcc cactaccggt actaggcaca 300

cggcttgtag ctnactggg cggttgtgtg ctatgaagtt cgctacactg gcctgtcaat 360

tatcttttgc atgcatatgc attatcatat acccaagtcg cgtacag 407

Sequence

<210> 1488

<211> 300

<212> nucleic acid

<213> Zea mays

<400> 1488

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tcgagattga gctggccgag gtcgaaatgc ccggcctcat ggcgtagcgc gccgagttcg 180

gcccgccaa gcccttcgcc ggcgctagga tctcggggtc tctccacatg accatccaga 240

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<210> 1489

<211> 259

<212> nucleic acid

<213> Zea mays

<400> 1489

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ccctccaggc tetgatggag ggtcttcagg tcttccctt ggaggacgtt gtctcggaag 120

ctgacatctt cgtgaccacc actggcaaca aggatatcat catggttgac cacatgagga 180

agatgaagaa caatgccatt gtctgcaaca ttggccactt tgacaatgaa attgatatgc 240

tggccttga gacctacc 259

<210> 1490  
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 <212> nucleic acid  
 <213> Zea mays  
 <400> 1490  
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 atcgacccca tctgcgccct ccaggctctg atggagggtc ttcaggtcct tcccttgag 180  
 gacgttgtct cggaagctga catcttcgtg accaccactg gcaacaagga tatcatcatg 240  
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 aat 303

<210> 1491  
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 <212> nucleic acid  
 <213> Zea mays  
 <400> 1491  
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 cgtgctcacc atcatccgcg acgggctcaa ggctgacccc aagaagtacc gcaagatgaa 180  
 ggagangctt gtcggcgtct ttgaggagac caccangggg gtcaagaggt ctaccagatg 240  
 caggagaccg gcgccctcct cttccctg 268

<210> 1492  
 <211> 278  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1492  
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 ttgacaacct gtatggttgc cgccactcgc tccctgatgg tctgatgagg gccactgacg 180  
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ctgctgccct caagcaggct ggtgccgtgt catgtgac 278

<210> 1493  
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 <212> nucleic acid  
 <213> Zea mays

<400> 1493

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 ccttnggtgt gnaactggcc atcctagctt tgtcangtnc tgctcattca ctaaccaggt 180  
 cattgcccaa cttgaactgt ggaaggagaa gagctctggc aagtatgaga agaagggtga 240  
 tgtgctcccc aagcaccttg atgagaaggt tgctganctc ca 282

<210> 1494  
 <211> 305  
 <212> nucleic acid  
 <213> Zea mays

<400> 1494

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 atggcggtgc gcgcgcaggt cgccccgtcc aagcccttcg ccggcgctag gatctcgggg 180  
 tctctccaca tgaccatcca gaccgccgtc ctcatcgaga cctcaccgc gctcggcgcc 240  
 gaggtccgtg gtgtcctgca acatttctcc acnaggacca gccgcgcgca tgcgcggaan 300  
 ggcg 305

<210> 1495  
 <211> 284  
 <212> nucleic acid  
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<400> 1495

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 agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180

cgccaagcc cttcgccggc gctaggatct cgggggtctct ccacatgacc atccagaccg 240  
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<210> 1496  
<211> 263  
<212> nucleic acid  
<213> Zea mays  
  
<400> 1496

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gcaggagacc ggcgcctcc tcttcctgc cattaacgtc aacgattccg tcaccaagag 120  
caagtttgac aacctgtatg gttgccgcca ctgcgtccct gatggtctga tgagggccac 180  
tgacgttatg atcgccggaa aggttgccgt ggtctgcgga taccgtgatg tcggcaaagg 240  
gttgtgctgc tgccctcaag cag 263

<210> 1497  
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<212> nucleic acid  
<213> Zea mays  
  
<400> 1497

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gcctcatggc gtgccgcgcc gagttcggcc cgtccaagcc cttcgccggc gctaggatct 240  
cgggggtctct ccacatgacc atccagaccg ccgtcctcat cgagaccctc accgcgctcg 300  
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<210> 1498  
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<212> nucleic acid  
<213> Zea mays  
  
<400> 1498

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agtatgagaa gaagggtgtat gtgtctccca agcaccttga tgagaagggt gctgctctcc 180  
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<210> 1499  
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 <212> nucleic acid  
 <213> Zea mays

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aaggtgtatg tgctcccaaa gcaccttgat gagaagggtg ctgctctcca cttgggcaag 240  
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ccgcngtntt cgcttaaang gggaaaccct cgngnat 277

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gaaattgata tgctcggcct tgagacctac cctggcgctca agcgcatcac catcaagccc 180  
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gttccccgag accaactg gcatcattgt ccttgctgag gtcgcctga tgaaccttgg 180  
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 gtgccgcgcc gagttcggcc cgtccaagcc cttcgcgggc gctaggatct cggggtctct 240  
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 aatgaaattg atatgctcgg ccttgagacc taccctggcg tcaagcgcat caccatc 237

<210> 1516  
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gatgaaggag aggtttgtcg gcgtctctga ggagaccacc acgggtgtca agaggctcta 180  
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 catggcggtgc cgcgcgcgagt tcggcccgtc caancccttc gccggcncta ggatntcggg 240  
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 gctcggcctt gagacctanc ctggcgctcaa ggcacacca tcaagcccca gactgancgc 180  
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 atccacgagg gtgtcaaggc cgaggaggat tacgagaaga ccggcaagat tccgacccgg 180  
 agtccaccga caacgttgag ttcaagatcg tgctcaccat catccgcgac gggctcaagg 240  
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 atgaaattga tatgctcggc cttgagacct accctggcgt caagcgcac accatcaagc 180  
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 tcatggcgtg ccgcgcgcgag ttcgggcccg ccaagccctt cgccggcgct aggatctcgg 240  
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 cccgtccaag cccttcgccg gcgctaggat ctcggggtct ctccacatga ncatccagac 240  
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 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180  
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 tctctcgcag gggacttcg gccgcctcga gattgagctg gccgaggtcg aaatgcccg 180  
 cctcatggcg tgccgcgccg agttcggccc gtccaagccc ttcgccggcg ctaggatctc 240  
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<210> 1547  
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 <400> 1547  
  
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 ctcgagattg agctggccga ggtcgaaatg cccggcctca tggcgtgccg cgccgagttc 180  
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 tacgagaaga ccggcaagat ccccgacccg gagtcancgg acaacgctga gttcaagatc 180  
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 ttggaggacg ttgtctcgga agctgacatc ttcgtgacca ccaactggcaa caaggatatc 180  
 atcatggttg accacatgag gaagatgaag aacaatgcc a ttgtctgcaa cattggccat 240  
 ttgacaatga attgatatgc tcggccttga gacctac 277

<210> 1550

<211> 277  
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 agaccggcna gatccccgac ccgngtnca ccgacaacgc tgagttcaag atcggtctca 180  
 ccatcntccg ngacgggctn aacgtgacc ccaagaagta ccgcnaantg aangagangt 240  
 tgtaggcgtc totgangaga ncacnacggg tgtnaag 277

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 <212> nucleic acid  
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 gggttgtgct ctccacttgg gcaagcttgg tgccaagctg accaagctca ccaagtctca 180  
 ggccgactac atcagcgtgc cgatcgaggg tccttacaag cctgcccact accggtacta 240  
 ggcacacggc ttgcagctca ctcgggccgt tgtgtgctat gaagttcgct a 291

<210> 1552  
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 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180  
 gcgtgccgcg ccgagttcgg ccggtccaag cccttcgccg gcgctaggat ctcggggtct 240  
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<210> 1553



<211> 318  
 <212> nucleic acid  
 <213> Zea mays  
  
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 gaaggttgct gctctccact tgggcaagct tggtgccaag ctgaccaagc tcaccaagtc 180  
 tcaggccgac tacatcagcg tgccgatcga gggtccttaa caagcctgcc cactaccggt 240  
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 cngggcctgn aatnattt 318

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 agaagggtgta tgtgctcccc aagcancttg atgagaaggt tgctgctctc cacttgggca 180  
 agcttgggtgc caagctgacc aagttcacca agtctcaggc cg 222

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 <213> Zea mays  
  
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<210> 1556

<211> 260  
 <212> nucleic acid  
 <213> Zea mays  
  
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 cggacttcgg ccgcctcgag attgagctgg ccgaggtcga aatgcccggc ctcatggcgt 180  
 gccgcgccga gttcggccccg tccaagccct tcgccggcgc taggatctcg gggctctctcc 240  
 acatgaccat ccagaccgcc 260

<210> 1557  
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 <212> nucleic acid  
 <213> Zea mays  
  
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 ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120  
 tctcgcaggc ggacttcggc ccgcctcgaga ttgagctggc cgaggtcgaa atgcccggcc 180  
 tcatggcgtg ccgcgcgcgag ttccggccccg ccaagccctt cgcgggcgct aggatctcgg 240  
 ggtctctcca catgaccatc cagaccgccc c 271

<210> 1558  
 <211> 223  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1558  
  
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 accatcaagc ccagactga ccgctgggtg ttccccgaga ccaacactgg catcattgtc 180  
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<210> 1559  
 <211> 293  
 <212> nucleic acid

<213> Zea mays

<400> 1559

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aggcggactt cggcgcgcct gagattgagc tggccgaggt cgaaatgccc ggccatcatgg 180  
cgtgccgcgc cgagttcggc cgtccaagc cttcgccgg cgctaggatc tcggggtctc 240  
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<210> 1560

<211> 244

<212> nucleic acid

<213> Zea mays

<400> 1560

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agattgagct ggccgaggtc gaaatgccc gctcatggc gtgccgcgcc gagttcggcc 180  
cgtccaagcc cttcgccggc gctaggatct cggggtctct ccacatgacc atccagaccg 240  
cogt 244

<210> 1561

<211> 358

<212> nucleic acid

<213> Zea mays

<400> 1561

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caaggtggtg ttnttgccct caagaaggcn gtgcccgtgt canttgaccg agatcgaccc 180  
catctgtgcc tccaggctct gatggagggt cttcaggctt tcccttgag gacgttgtct 240  
ctgangtgac atcttcgtga ccaccactgg caaccaagga tatcncatgg ttgancacat 300  
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<210> 1562

<211> 218  
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<400> 1562

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 gctgctctcc acttggggcaa gcttggtgcc aagctgacca agctcaccaa gtctcaggcc 180  
 gactacatca gcgtgccgat cgagggtccc tacaagcc 218

<210> 1563  
 <211> 269  
 <212> nucleic acid  
 <213> Zea mays

<400> 1563

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 gatctctcgc aggcggactt cggccgcctc gagattgagc tggccgaggt cgaaatgcc 180  
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<400> 1564

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 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggccctcatg 180  
 gcgtgccgcg ccgagttcgg ccgtccaag cccttcgccg gcgctaggat ctcggggtct 240  
 ctccacatga ccatccagac 260

<210> 1565  
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 <212> nucleic acid

<213> Zea mays

<400> 1565

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cgatccggag tccaccgaca acgctgagtt caagatcgtg ctcaccatca tccgcgacgg 180  
gctcaaggct gacccaaga agtaccgcaa gatgaaggag aggttgtcgg cntctctgag 240  
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<210> 1566

<211> 282

<212> nucleic acid

<213> Zea mays

<400> 1566

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tgggcaagct tggtgccaag ctgaccaagc tcaccaagtc tcaggccgac tacatcagcg 180  
tgccgatcga ggggtccctac aagcctgccc actaccggta ctaggcagcc agcacacggc 240  
ttgcagctca ctggggccgt tgtgtgctat gaagttcgct ac 282

<210> 1567

<211> 235

<212> nucleic acid

<213> Zea mays

<400> 1567

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gggtctcncc acatgaccat ccagaccgcc gtctcctcgc agacntcac ngcgtcggc 180  
gccgaggtcc gtggtgcnc ngcaacatct tctccagcag gancacgccg ccgcc 235

<210> 1568

<211> 239

<212> nucleic acid

<213> Zea mays

<400> 1568

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 gaccatccag accgccgtcc tcatcgagac cctcaccgcg ctcggcgccg aggtcngctg 180  
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<210> 1569

<211> 250

<212> nucleic acid

<213> Zea mays

<400> 1569

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 ggacttcggc cgcctcgaga ttgagctggc cgaggtcgaa atgcccggcc tcattggcgtg 180  
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<210> 1570

<211> 264

<212> nucleic acid

<213> Zea mays

<400> 1570

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 gctcatggc gtgccgcgcc gagttcggcc cgtccaagcc cttcgccggc gctaggatct 240  
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<210> 1571

<211> 274

<212> nucleic acid

<213> Zea mays

<400> 1571

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 gggcaagctt ggtgccaagc tgaccaagct caccaagtct caggccgact acatcagcgt 180  
 gccgatcgag ggtccctaca agcctgcccc ctaccggtac taggcnacgg cttgcagctt 240  
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<210> 1572  
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 cttcggccgc ctgagattg agctggccga ggtcgaaatg cccggnetca tggcgtgccg 180  
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 <212> nucleic acid  
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<400> 1573

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 ttcttcatct tcctcatgtg gtcaaccatg atgatatcct tgttgccagt ggtggtcacg 180  
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<210> 1574  
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 <213> Zea mays

<400> 1574

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 cctcatggcg tgccgcgcg agttcggccc gtccaagcnc ttcgccggcg taggattcgg 240  
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<210> 1575  
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 <212> nucleic acid  
 <213> Zea mays

<400> 1575  
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 cgctcgaga ttgagctggc cgaggtcgaa atgcccgcc tcattggcgtg ccgcgccgag 180  
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<210> 1576  
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<400> 1576  
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 ggttgtctgt ctccacttgg gcaagcttgg tgccaagctg accaagctca ccaagtctca 180  
 ggccgactac atnccgcgtg ccgatcgagg gtccctacaa gcctgccac ttaccggtat 240  
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<210> 1577  
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caagggtcaag gatctctcgc aggcggactt cggccgcctc gagattgagc tggccnaggt 180  
cgaaatgccc ggcctcatgg cgtgccgcgc cgagttcggc ccgtccaagc ccttcgccgg 240  
cgctaggatc tcgggggtctc tccacatgac ca 272

<210> 1578

<211> 179

<212> nucleic acid

<213> Zea mays

<400> 1578

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gtttgacaac ctgtatggtt gccgccactc actccctgat ggtctgatga gggccaccga 120  
cgttatgac gccggtaagg ttgccgtggt ctgccgatac ggtgatgttg gcaagggtt 179

<210> 1579

<211> 278

<212> nucleic acid

<213> Zea mays

<400> 1579

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tggaagaagac ctggtctgga cgggagtaca aggtcaagga tctctcgag gcggacttcg 120  
gcgcctcga gattgagctg gccgaggtcg aaatgcccg cctcatggcg tgccgcgcgc 180  
agttcgcccc gtccaanccc ttngccggcg taggatctcg ggtcctcca catgaccatc 240  
cagaccgcng tentatcgag acctnacn gn gttggngg 278

<210> 1580

<211> 243

<212> nucleic acid

<213> Zea mays

<400> 1580

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gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgcaggc ggacttcggc 120  
cgctcgaga ttgagctggc cgaggtcgaa atgcccggcc tcatggcgtg ccgcgccgag 180  
ttcggcccgt ccaagccttt cgccggcgta ggatctcggg gtctctccac atgaccatcc 240  
aga 243

<210> 1581  
<211> 247  
<212> nucleic acid  
<213> Zea mays  
<400> 1581

ctcgttccat ttcccattct ccagatcca attcgcgagt tctccctcct ctgcggccat 60  
ggcgtctctt gtggagaaga cctcgtctgg acgggagtac aaggtaagg atctctcgca 120  
ggcggacttc ggcgcctcg agattgagct ggccgaggtc gaaatgcccg gcctcatggc 180  
gtgcgcgcgc gagttcggcc cgtccaagcc cttcgccggc gctaggatct cggggtctct 240  
ccacatg 247

<210> 1582  
<211> 246  
<212> nucleic acid  
<213> Zea mays  
<400> 1582

cgttccatt tcccattctc ccagatccaa ttcgcgagtt ctccctcctc tgccgccatg 60  
gogctctctg tggagaagac ctcgtctgga cgggagtaca aggtcaagga tctctcgcag 120  
goggacttcg gcgcctcga gattgagctg gccgaggtcg aaatgcccg cctcatggcg 180  
tgccgcgcgc agttcggccc gtccaagccc ttcgccggcg ctaggatctc ggggtctctc 240  
cacatg 246

<210> 1583  
<211> 276  
<212> nucleic acid  
<213> Zea mays  
<400> 1583

cctccctcac tctcgttcca tttcccattc tcccagatcc aattcgcgag ttctccctcc 60

tctgcggccca tggcgctctc tgtggagaag acctcgtctg gacgggagta caaggtcaag 120  
gatctctcgc agggcgactt cggccgcctc gagantgagc cggccgaggg cgaaaanccc 180  
ggcctcaagg cgtgccgcgc cgagtnggcc cgnccaagcc cttcgccggc gctaggatct 240  
cggggtctct ccacatgacc atccagaccg ccgtcc 276

<210> 1584  
<211> 178  
<212> nucleic acid  
<213> Zea mays

<400> 1584

attgtotgca acattggcca ctttgacaat gaaattgata tgctcggcct tgagacctac 60  
cctggcgtea agcgcatcac catcaagccc cagactgacc gctgggtgtt ccccagagacc 120  
aacactggca tcattgtcct tgctgagggc cgctganga nccttggtgtg tactnntg 178

<210> 1585  
<211> 175  
<212> nucleic acid  
<213> Zea mays

<400> 1585

cgacggcggc gacgccacgc tgctcatcca cgaggggtgtc aaggccgaga aggagtacga 60  
gangaccggc aagatccccg acccgagtc caccgacaac gctgagttca agatcgtgct 120  
caccatcatc cgcgacgggc tcaaggctga cccaagaag taccgcaaga tgaag 175

<210> 1586  
<211> 235  
<212> nucleic acid  
<213> Zea mays

<400> 1586

tcccatctc ccagatccaa ttcgcgagtt ctcctctc tgccgccatg gcgctctctg 60  
tggaagaagac ctgctctgga cgggagtaca aggtcaagga tctctcgag gcggacttcg 120  
gcgcctcga gantgagctg gccgaggtcg aaatgcccg cctcatggcg tgccgcgccg 180  
agttcgcccc gtccaagccc ttcgccggcg ctaggatctc ggggtctctc cacat 235

<210> 1587

<211> 180  
 <212> nucleic acid  
 <213> Zea mays

<400> 1587

ancaagtttg acaacctgta tggttgccgc nactcgctcc ctgatggtct gatgagggcc 60  
 actgacgtta tgatcgccgg aaaggttgcc gtggtctgcy gatacgggtga tgtcgnnang 120  
 ggttggtgng ctgcnctcaa gcaggctggt gcccggtgna tctgtancgn gatcgacccc 180

<210> 1588  
 <211> 236  
 <212> nucleic acid  
 <213> Zea mays

<400> 1588

cccgttccat ttcccatct cccagatcca attcgcgagt tctccctcct ctgccgccat 60  
 ggcgctctct gtggagaaga cctcgtctgg acgggagtag aaggtcaagg atctctcgca 120  
 ggcggacttc ggccgcctcg agattgagct ggccgaggte gaaatgcccg gcctcatggc 180  
 gtgccgcgcc gaggtcggcc cgtccaagcc cttcgccggc gctaggatct cgggggt 236

<210> 1589  
 <211> 272  
 <212> nucleic acid  
 <213> Zea mays

<400> 1589

ccttgggagg acgttgtctc ggaagctgac atcttcgtga ccaccactgg caacaaggat 60  
 atcatcatgg ttgaccacat gaggaagatg aagaacaatg ccattgtctg caacattggc 120  
 cacattgaca aagaaatgnt angcncgggc ccgnagaccn aacccggcgg caanngnanc 180  
 aacaacnagg cccagacgga ncgcccgggg gnccccgaga ccaacacggn aacaagtcct 240  
 gncgaggggc gcccgangaa ccatngggga gc 272

<210> 1590  
 <211> 260  
 <212> nucleic acid  
 <213> Zea mays

<400> 1590

ccctcactcc cgttccattt ccccatctcc cagatccaat tcgcgagttc tccctcctct 60  
 gccgccatgg cgtctctgtt ggagaagacc tcgtctggac gggagtacaa ggtcaaggat 120  
 ctctgcagg cggacttcgg ccgcctcgag attgagctgg ccgaggtcga aatgcccggc 180  
 ctcatggcgt gcgcgcgga gttcggcccg tccaagcent tcgccggcgc taggatctcg 240  
 ggggtctctc cacatganca 260

<210> 1591  
 <211> 245  
 <212> nucleic acid  
 <213> Zea mays

<400> 1591

cctcaactccc gttccttttc cccntctccc agntccantt cgcgagttct cctcctctg 60  
 ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggntc 120  
 tctgcaggc ggacttcggc ccgcctcgaga ttgagctggc cgaggtcga atgcccggcc 180  
 tcatggcgtg ccgcgcgag ttcggcccg ccaagccctt cgcggcgct aggatctcg 240  
 ggtct 245

<210> 1592  
 <211> 209  
 <212> nucleic acid  
 <213> Zea mays

<400> 1592

cccagatcca attcgcgagt nctccctcct ctgccgccat ggcgctctct gtggagaaga 60  
 cctcgtctgg acgggagtac aaggtcaagg atctctcgca ggcggacttc ggccgcctcg 120  
 agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180  
 cgtccaagcc cttcgcgggc gctaggatt 209

<210> 1593  
 <211> 221  
 <212> nucleic acid  
 <213> Zea mays

<400> 1593

cccatctccc agatccaatt cgcgagttct cctcctctg ccgccatggc gctctctgtg 60

gagaagacct cgtctggacg ggagtacaag gtcaaggatc tctcgcaggc ggacttcggc 120  
 cgcctcgaga ttgagctggc cgaggtcgaa atgcccggcc tcatggcgtg ccgcgccgag 180  
 ttcggccccgt cnaagccctt cgcnggcggt aaggntttgg g 221

<210> 1594  
 <211> 226  
 <212> nucleic acid  
 <213> Zea mays

<400> 1594

ctctcgttcc atttcccat ctcccagatc caattcgcga gttctccctc ctctcgggcc 60  
 atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa ggatctctcg 120  
 caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180  
 gcgtgccgcg ccgagttcgg ccggtccaag cccttcgccg gcgcta 226

<210> 1595  
 <211> 149  
 <212> nucleic acid  
 <213> Zea mays

<400> 1595

ccctcgagga gtactggtgg tgcaccgagc gctgcctcga ctggggcgag gcggggcgcc 60  
 ccgacctcat cgtcgacgac ggcgggcgag ccacgctgct catccacgag ggtgtcaagg 120  
 ccgaggagga ttacgagaag accggcaag 149

<210> 1596  
 <211> 301  
 <212> nucleic acid  
 <213> Zea mays

<400> 1596

cccgtncat ntccccatn ncccagatcc aatttcgcga gtacnaccnc ctcnagncgc 60  
 catggggggt ctctgtggag aagacatcgt ctggacggga gtacaagggtc aaggatcnct 120  
 cgcaggcgga cntcgccgc ntcgagattg agcnggccga ggtcgaaatn acccggnctc 180  
 atggcgtnen gcgcgagtt cggcccgtcc aagcncttcg ccggcgctan gattcgggggt 240  
 ctctaccaca tgaccatcca gacggcgctc tcatcgagac atcacagcgt cggngccgag 300

t

301

<210> 1597  
 <211> 174  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1597

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 tccccgaccc ngngtccacc gataacnctg nnttcaagat cgtggctcac catcatccgc 120  
 gacgggctca aggctgacnc caagaagtac cgcaagatga aggagnggct ngtt 174

<210> 1598  
 <211> 228  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1598

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 gcgcgcatgg cgctctctgt ggagaagacc tcgtctggac gggagtacaa ggtcaaggat 120  
 ctctcgcagg cggacttcgg ccgcctcgag attgagctgg ccgaggtcga aatgcccggn 180  
 ctcatggcgt gccgcgccga gttcggcccg tccaagccct tcgccggg 228

<210> 1599  
 <211> 227  
 <212> nucleic acid  
 <213> Zea mays  
 <400> 1599

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 cctcgtctgg acgggggtac aaggtcaagg atctctcgca ggcggacttc ggccgcctcg 120  
 agattgagct ggccgaggtc gaaatgcccg gcctcatggc gtgccgcgcc gagttcggcc 180  
 cgtccaagcc cttcgccggc gtaggatctc ggggtcctca natgaca 227

<210> 1600  
 <211> 236  
 <212> nucleic acid  
 <213> Zea mays

<400> 1600

cggttccatt tccccatctc ccagatccaa ttcgcgagtt ctccctcctc tgccgccatg 60  
gcgctctctg tggagaagac ctcgctctgga cgggagtaca aggtcaagga tctctcgcag 120  
gcggaacttcg gccgcctcga gattgagctg gccgaggtcg aaatgcccgg cctcatggcg 180  
tgccgcgcgcg agttcggccc gtccaagcct tcgccggcgc taggatctcg gggttc 236

<210> 1601

<211> 209

<212> nucleic acid

<213> Zea mays

<400> 1601

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atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaaggtaa ggatctctcg 120  
caggcggact tcggccgcct cgagattgag ctggccgagg tcgaaatgcc cggcctcatg 180  
gcgtgccgcg ccgagttcgg cccgtccaa 209

<210> 1602

<211> 426

<212> nucleic acid

<213> Zea mays

<400> 1602

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ccgccanggn gctctctgtg gagaagacct cntctggacg ggagtacaag gtcaaggatn 120  
nctcgcaggn ggacttcggt cggcncgaga ttganctggc ngagncgaa atgcccggcc 180  
ncatggcntg ccgngccgan ttcggcccgt ctangccctt cgnccgcgct aggnnacng 240  
ngtctacca catgaccatc naagactgtn ttcctcancg agaactnang gtacgacggt 300  
accnaagtnc ccngatatcc gngnncnact tcccaacant tgaggaaact acantgcatt 360  
tcnanngacc gnaccanaat tgncanatca aaggtaaatt aanncntagn ctncaanggg 420  
naantg 426

<210> 1603

<211> 232

<212> nucleic acid



<213> Zea mays

<400> 1603

ccccatctcc cagatccaat tcgcganttc tccctcctct gcggccatgg cgctctctgt 60  
ggagaagacc tcgtctggac gggagtagaa ggtcaangat ctctcgcagg cggacttcgg 120  
ccgntcnat attgagctgg ccgaggtcga aatgcncggc ctcatagcgt gccgcgccga 180  
tttcggncog tcnagccct tcgcgggng ctaggatctc ggggtctctc ca 232

<210> 1604

<211> 218

<212> nucleic acid

<213> Zea mays

<400> 1604

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ctgcgcgcac ggcgtctctt gtggagaaga cctcgtctgg acgggagtag aaggtaagg 120  
atcnctcgca ggccgaentc ggccgcctcg agantgagcn ggccgaggtc gaaatgcccg 180  
gcctcatggc gtgcgcgcgc gagttcggcc cgtccaag 218

<210> 1605

<211> 134

<212> nucleic acid

<213> Zea mays

<400> 1605

gccgcacac actccctgat ggtctgatga gggccaccga cgttatgatc gccggtagg 60  
ttgccgtggt ctgcggatac ggtgatgttg gcaagggttg tgccgctgca ctcaagcagg 120  
ctggtgcccg tgtc 134

<210> 1606

<211> 152

<212> nucleic acid

<213> Zea mays

<400> 1606

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catcttctcc acgcaggacc acgccgccgc cgcnatcgcg cgcgantcgg ccgccgtgtt 120

cgctgnaaa gggggagacc cttgaggagt ag

152

<210> 1607  
<211> 287  
<212> nucleic acid  
<213> Zea mays

<400> 1607

gtccaagtat gagaagaagg tgtatgtgct cccaagcac cttgatgaga aggntgctgc 60  
nnctcccact tgggcaagct tggtgccaag ctgaccaagc tcaccaagtc tcaggccgac 120  
tacatcagcg tgccgatcga gggtccttac aagcctgccc actaccggtg ctaggcagcc 180  
agcacacggc ttgcagctca cttcggggccg ttgtgtgcta tnaagtnenc ncncaactgnc 240  
ctgtcagttc atcttttgca tgcatatgca ntatcatata cgcacg 287

<210> 1608  
<211> 123  
<212> nucleic acid  
<213> Zea mays

<400> 1608

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ccacatgagg aagatgaaga acaatgccat tgtctgcaac attggccact ttgacaatga 120  
ant 123

<210> 1609  
<211> 348  
<212> nucleic acid  
<213> Zea mays

<400> 1609

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gccctcaagc aggttggtgc ccgtgtcatt gtgaccgaga tcgaccccat ctgtgcnctc 120  
caggctctga tggagggctt caggtcttcc cttggaggac gtgtctctga agcnnacatc 180  
tcgtgacaac attggcanca agtatcatca tggtgaccac atgaggaaga gaagacccat 240  
gccatgtctg cacattggca ctttgacatg aattgatatc tcggcttgag actacctgng 300  
tcaangctca catcagcccn gatgacgtgg tgtcccgaga canatgga 348

<210> 1610  
 <211> 266  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1610  
  
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 tcaaggccga gcatatcaat ttccattgtc aaagtggcca atgttgcaga caatggccag 120  
 tgggttticat ccttctcat gtggtcaacc atgatgaata tccttgttgc cagtgggtgg 180  
 tcaogaagat gtcagctttc cggaggnaaa cggctccann ctgtcgtga gaanncgtgt 240  
 tgengnaggg gtgcccgtgc cgngc 266

<210> 1611  
 <211> 143  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1611  
  
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 tcattgtgac cgagatcnac cccatctgtg cctccaggc tctgatggag ggtcttcagg 120  
 tcnttccctt ggagnacgtt ntt 143

<210> 1612  
 <211> 118  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1612  
  
 aacgattccg tcaccaagag caagtttgac aacctgtatg gttgccgcca ctactccct 60  
 gatggtctga tgagggccac cgacgttatg atcgccggtg aggttgccgt ggtctgcg 118

<210> 1613  
 <211> 265  
 <212> nucleic acid  
 <213> Zea mays  
  
 <400> 1613  
  
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ctctgccgcc atggcgctct ctgtggagaa gacctcgtct ggacgggagt acaagggtcaa 120  
ggattctcgc aggcggactt cggccgcctc gagattgagt ggccgaggtc gaaatgccgg 180  
ctcatggcgt gccgcggcga gttcggtccc tccaagcctt cggcggcgta agatctcggg 240  
gtctttcaca tgaacatcag accgc 265

<210> 1614  
<211> 111  
<212> nucleic acid  
<213> Zea mays  
<400> 1614

catggttgac cacatgagga agatgaagaa caatgccatt gtctgcaaca ttggccactt 60  
tgacaatgaa attgatatgc tcggccttga gacctaccct ggcgtcaagc g 111

<210> 1615  
<211> 154  
<212> nucleic acid  
<213> Zea mays  
<400> 1615

cnccagatc caattcgcga gtnncnccnc ctctgccgcc atggcgctct ctgtggagaa 60  
gacntcgtct ggacgggagt acaagggtcaa ggatctctcg caggcggacn tcggccgcct 120  
cgagattgag cnggccgagg tcgaaatncc cggc 154

<210> 1616  
<211> 226  
<212> nucleic acid  
<213> Zea mays  
<400> 1616

cccagatcca attcgcgagt tctccctcct ctgccgccat ggcgctctct gtggagaaga 60  
cctcgtcctg gaacgggagn acaagggtcaa ggntntntng naggngnant tnggcngcnt 120  
cnaattnan ctggccnagg tcgaaatgcc cggcntnatg gcgtgcngcg ccganttcgg 180  
cccgccaag ccttcgccgg cgnaggntc tcgggggtctt cnacat 226

<210> 1617  
<211> 229  
<212> nucleic acid

<213> Zea mays

<400> 1617

ggatatcatc atggttgcca gtggtggtca cgaagatgtc agcttcagag acaacgtcct 60  
ccaagggaag gacctgaaga cctccatca gagcctggag ggcacagatg gggtcganct 120  
cggtcacaag gntatcatcn ngggtgacca catgaggag atgaagaaca atgccattgt 180  
ctgcaacatt ggccactttg acaatgaaat tgatatgtc ggccttgag 229

<210> 1618

<211> 120

<212> nucleic acid

<213> Zea mays

<400> 1618

ctttgtontg tctgtctcat tcactaacca ggtcattgcc caacttgaac tgtggangga 60  
gaagagctct ggcnagtatg anaagaaggt gtatgtgtc cccaagcacc ttgatgagaa 120

<210> 1619

<211> 109

<212> nucleic acid

<213> Zea mays

<400> 1619

gtgccctcca ggctctgatg gaggggtcttc aggtccttcc cttggaggac gttgtctctg 60  
aagctgacat cttcgtgacc accactggca acaaggatat catcatggt 109

<210> 1620

<211> 96

<212> nucleic acid

<213> Zea mays

<400> 1620

ggttgaccac atgaggaaga tgaagaacaa tgccattgtc tgcaacattg gccactttga 60  
caatgaaatt gatatgtctg gccttgagac ctaccc 96

<210> 1621

<211> 118

<212> nucleic acid

<213> Zea mays

<400> 1621

cgctgagttc aagatcgtgc tcaccatcat ccgcgacggg ctcnaggctg accccaagaa 60  
gtnccgcaag atgaaggaga ngcttgctcg cgtctctaag gagancancc angggtgt 118

<210> 1622

<211> 559

<212> nucleic acid

<213> Zea mays

<400> 1622

gnnnnnnngnn tnnonagann nnttcncttt ccaacnctc ttgaatttcc gggtcgaccc 60  
acgcgtccgc aacctgtatg gntgccgnc a ctngatccct gatggtctga tgagggccac 120  
tgaccgttat gntcgccgga aaggttgccg tggctcncgg atacngtgat ntcngcaagg 180  
gttgtgctgc tgcaaatnaa gcanggctng tgcccgtnct attgtnaccc gagancnacn 240  
natctgtncn nntacangct cttattngaa ggtctttang nccttcnctt ggaaganngt 300  
ggctntgaag ctncatnttn ngaccaccac tgnaaacaag gatatnnnat ggttgaccac 360  
atgangaana tgaagnacaa tgccattggc tgnaacattg ggccactttt gacaatgaan 420  
ttgatatgct cnggccttga gacctaccct ggcgtcaaag cgcattatnc atcaaanccc 480  
anactgaccg cttgggtggt tcengagacc aaacactggc atcattgggc cttnctgaag 540  
ggtcnnctgg ntinaacntt 559

<210> 1623

<211> 88

<212> nucleic acid

<213> Zea mays

<400> 1623

cgacaacgct gagttcaaga tcgtgctcac catcatccgc gacgggctca aggctgaccc 60  
caagaagtac cgcaagatga aggagagg 88

<210> 1624

<211> 82

<212> nucleic acid

<213> Zea mays

<400> 1624

atctttctcca cgcaggacca cgcgcgcgcc gccatcgcg cgcactcggc cgcogtggtc 60  
gcctggaagg gggagaccct cg 82

<210> 1625  
<211> 139  
<212> nucleic acid  
<213> Zea mays

<400> 1625

cctcactccc gttccatttc cccatctccc agatccaatt cgcgagttct cctcctctg 60  
ccgccatggc gctctctgtg gagaagacct cgtctggacg ggagtacaag gtcaaggatc 120  
tctcgcangc ggacttcgg 139

<210> 1626  
<211> 255  
<212> nucleic acid  
<213> Zea mays

<400> 1626

agcttggtgc caagctgacc aagctcacca agtctcaggc cgactacatc agcgtgccga 60  
tcgagggtcc ctacaagcct gccactacc ggtactaggc acacggcttg cagctcactc 120  
gggcccgttg gtgctatgaa gttcgctaca ctggcctgtc agttatcttt tgcattgcata 180  
tgcattatca tatacgcagt cgcgtanagg ttttcttatg gttatcgctt gancngnngn 240  
gggagggaag gagct 255

<210> 1627  
<211> 224  
<212> nucleic acid  
<213> Zea mays

<400> 1627

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<210> 1628

<211> 113  
 <212> nucleic acid  
 <213> Zea mays

<400> 1628

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<210> 1629  
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 <212> nucleic acid  
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<400> 1629

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<210> 1630  
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 <212> nucleic acid  
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<400> 1630

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<210> 1631  
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 <212> nucleic acid  
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<400> 1631

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 gagtgcgatg tctacgtcaa gccagagctg ttcagcgctg gaggcagtgt taaggacaga 180  
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cttatcgagc ctaccagtgg aaacactggt atcgggtcttg ctc 283

<210>	1632
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<212>	nucleic acid
<213>	Zea mays

<400>	1632
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gtgtaggcgc tcggatgctt ttctacccga cgctgggtgta caacgtcgtg aggaatcggc 180

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catccctagc gatgttctcc ggctaaaga 269

<210>	1633
<211>	125
<212>	nucleic acid
<213>	Zea mays

<400>	1633
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ttgtggcctg aagacnccgt ccatttgacn ncatncnggc tgaagtnccg tncctncttc 120

gatgt 125

<210>	1634
<211>	123
<212>	nucleic acid
<213>	Zea mays

<400> 1634

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ccatcaaggc cccttggtggc ctgaagactc gtccattcga ctcaattctg gctgaagtgc 120

gcg 123

<210>	1635
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<212>	nucleic acid
<213>	Glycine max

<400> 1635

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gtgtgatcag atatctgatg cagtgcctga tgcctgcctg gagcaagacc ctgaaagcaa 180  
ggttgcctgt gagacatgca ccaagaccaa cttgggtcatg gtatttggag agatcaccac 240  
aaaagccaat atngactatg agaagattgt gcgcgatacg tgccgttcta taggatttgt 300  
gtctggtgat gt 312

<210> 1636

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 1636

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gcacggcaac ttaaccaaag acccgaggaa atcgggtgtg agaccaggggt cactgtttgc 180  
tatgcacgga cgagacccca gattgngcca tgagtcattgt ncttncaata aactcgggtgt 240  
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<210> 1637

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 1637

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acagcatgat gagnctgtc acaaatgatg ngattgcagc tganctnaaa gaacacgtga 180  
ttaagcntgt gattccngng nagtaccttg nngagccgac cnntggccng tngaaccctc 240  
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<210> 1638

<211> 292

<212> nucleic acid

<213> Glycine max

<400> 1638

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acacgangan nontgtccnc cantgacgan attgctgntg nctcaaaga gcatgtgntc 240  
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<210> 1639

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 1639

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taaggccatt cattcctcac cgctgtgtgc tgggagtttt ttgagctttg ccottatcat 180  
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<210> 1640

<211> 275

<212> nucleic acid

<213> Glycine max

<400> 1640

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tgtaccttga gcagcaaagt cctgacatag cacaaggagt ccatggccac ctcaccaagc 180  
gaccagaaga cataggagct ggcgatcagg gtcacatgtt tggctacgcc acagatgaaa 240  
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<210> 1641

<211> 317

<212> nucleic acid

<213> Glycine max

<400> 1641

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ncttgnntgn gaagnccatt ttccagttca acccctctgg ccgttttgtc attngaggtc 180  
ctcaaongtg atgctctctc caccagccga caagatccat tcnnccgnta cttanggagg 240  
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<210> 1642

<211> 278

<212> nucleic acid

<213> Glycine max

<400> 1642

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gaagaaaacgc agaaactcat aaaccatcgc tgactccgca attttcgtcg gttcctcaca 180  
cactctctga tttaancncg gtgccaagct caccgaggtt cggaagaacg ggacatgccc 240  
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<210> 1643

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<212> nucleic acid

<213> Glycine max

<400> 1643

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tttttttagcg ttgcccttat aatgtctntt atccataact ttccacgtcc cttgctctgt 180  
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acatgatcaa ctaaaaaat 259

<210> 1644

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 agcctngtgc t 191

<210> 1645  
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 catatgacca agaaacctga ggaaattggg gctggtgacc aaggccacat gtttggttat 180  
 gctacagacg aaacacctga gttaatgcc ctcactcatg tgcttgctac taaacttnt 240  
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 acccaagtga cattgag 317

<210> 1646  
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 acccaaccaa ggttgatagg agtggtgcat acattgttag gcaagcagcc aagagcgtgg 180  
 tagcttcagg gcttgacga cgttgcatg tgcaggtttc ttatgcaatt ggagttccag 240  
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 tcccacaanc tgaactcggc catgaccaag gctcgcnagg attgtactct gccatggctg 180  
 cgacccgaca ccaagactca ggtcactgtc gagtacgcc acgatggcgg tgccgtcgtc 240  
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 aagctccgcg aggagatcaa ggagaagatc at 332

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 ccaaaagacc cgangnaata tcaggntctt gctgctcaag gtccttgtaa acattgagca 120  
 gcagagccct gatattgcc aggggtgtgca cggccacett accaaaagac ccgaggaaat 180  
 cgggtgctgga gaccagggtc acatgttttg ctatgccact gtcgagaccc cagaattgat 240  
 gccattgagt catgttcttg caactaaact cgggtgctcgt ctcaccgagg ttcgcaa 297

<210> 1649  
 <211> 348  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1649  
  
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 tgntnaagct anccatctct ntctctctnt cttaagnngn tccttngnna canntagaa 120  
 tggcctnaga aattncctat tcacnctnna tcagtgaacg ngnggcanc cgcacaggnn 180  
 ctgtgaccag atctccgatg ctgtgntcng attcatgctt ggagcaggac cctgagagca 240

aggttgctg tgaagcctgc aacaagacca acatggtggt ggttttcgga gagatcacia 300  
ccaaggccaa cgtggnctat ganaagattg tgcgggacac atgcaagg 348

<210> 1650  
<211> 324  
<212> nucleic acid  
<213> Glycine max

<400> 1650

ntcgcgtgca gcgtacgtaa gctcgggaatt cggctcgagg cgagtgttct ttcttcgttt 60  
caacacctin ntttgcannn gntgcttctt ctgcttgag anntggcaca agaaacnntt 120  
ctatncacat ctgantctgt anacgagggg caccocggac aagctgtgcy nccagatctc 180  
tgatggcagt gctcgatgcy tgctgnacn ggacctgag cagcgagggt gcctgtnaga 240  
catgcaccaa gacngnntg gncatggtct ttgngagat cagnanccaa ggncagcagt 300  
aggtatgag aannttnnnc gtga 324

<210> 1651  
<211> 318  
<212> nucleic acid  
<213> Glycine max

<400> 1651

tcacangcac gcgtacgtaa gctcgggaatt cggctcgagg cagacttcac ctgggaagtt 60  
gtgaagccac tcaagtcaga gaagcctcaa gcttaagagt gttgttaagt taatcactac 120  
ccttcagtgg ntgtcttgct ggggtgtggat gaataatttg cgtgtttcat gactactact 180  
actactactc cnttcnntgt ctaatgccat ctcatcnatn nctaaactgn tcgntttntt 240  
tttntccont atactcncaa tttgttggtt ggcnatgnaa tgtcactgtg ttgatgcatg 300  
gaattttatc caaangaa 318

<210> 1652  
<211> 312  
<212> nucleic acid  
<213> Glycine max

<400> 1652

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acaggaccca gacagcaagg ttgctgcga aacatgcacn aaaaccaact tggcatggn 120  
 cttcggagaa atcacgacca aggccaatgt tgactacgag aagatagtgc gtgtacaccc 180  
 tgagcaatat cagggctctg ctgctcaatg ttgacgagga ccttgagtt gtcggcatcc 240  
 agtcccacat cgtttgagac aaagccgatg ttcccgaata tcttggtgtt tggagcgatc 300  
 cgagaaaaaac tc 312

<210> 1653  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1653

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 nccgaggggc accccgacaa ncntgtgcga cnagatctct gatgcagtgc tcgatgcgtg 120  
 ccttgaacag gaccctgaca gcaagggttg ctgtgagaca tgcaccaaga ccaacatggt 180  
 catgggtcttt ggagagatca caaccagggc aangtanann ntgagaagan tgtccnggan 240  
 anangccgcg aaattggatt cnaccccgga nnanttggt nnnnnnnna aaantnnngg 300  
 ntnggncaan nnttgnnnc 320

<210> 1654  
 <211> 506  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1654

ggntnnnnnn aactnttacg ctctcaggtg cccgtcanag aattnacggg tcgagccacg 60  
 cgtcagtacg gatgcgagaa gacgacagaa gggggcagcg cttgagacca agccccactc 120  
 aaccaacaca acactctctc tgctctctct ctaactttca aagtttttta agtnnttaga 180  
 tggcagagac attcctaatt acctcaaagt cncnaacga agganacctg acaagctctg 240  
 cgaacaaatn tccgatgctg tncnnaanc ctgccttgaa caagacccaa acagcaaagt 300  
 tncctgccaa acatgcccc aagaccaactt tgcenangtc ctccgagaaa ttncaccaa 360  
 gggcaacntt nactacgana anantntgcg ttacacctgc nggaacatcg ggttcntctc 420  
 aaacgaagtt ggactttatg cctgacaaat gcaanggcc tnttaacaat ngncancaa 480



aaccctgaaa ntnnccaagg gggttg

506

<210> 1655  
<211> 501  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1655

ggnnnnnnnn nnnancttta actctccgcg ttcaggtaaa ggtttagaat tcccgggncg 60  
accacgcgt cnanccang cgtccgtacg gctgcgagag gangacagaa gggggcagcg 120  
cttgagacca agccccactc aaccaccaca ccaactctctc tgctcttctt ctatctttca 180  
acgttttttna agtattaaga tggcataaac attcctatatt acttcatant cagtnaanga 240  
gggacacnct gacgtinctc gcgannanatt ctengattct gtcctcnacn cntgccttna 300  
tnatgacnca cncancatcg ttgcttgnta natatacncc angaccaact tngtctnggt 360  
cttcgggagag atnnccatna tggcnaacgt tgactancat gaagatcntg cntnacacct 420  
gcaagaattc ggattcgtct caaaccatgt tggganntat gctgannntg naaggttcnt 480  
tgnaaacggt tgnccntttc a 501

<210> 1656  
<211> 533  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1656

angggtgata ttntgtgten aaaggatatag tangtgntag attntcttcg gtttcggtat 60  
taccnacnng accacgcgt cccgcccacg cgtccggctg cgataagacg acagaanggg 120  
atacctatgg tggctggggg gngggatggg ggaagtgcct ttggggggaa ngaccctacc 180  
aaggttgaca gaagtgggtc ctatattgta aggcattgctg caaanagtgt ngtggcaaat 240  
ggccttgcta naagggtgcat tgtgcaagtt tcctatncca ttggtgtccc tgaacccttg 300  
tcanatgttt ntngacactt atngaactgn naanattcna nacaannagag attttinctat 360  
anntatanga ataattttta nnttnananc tngnnnnngtn tnncataaan nttingtaant 420  
aaataagnnn naantntttt gtnnatttaa naaattntan tnttnnttta natagnnaaa 480  
taatttttna agtnatttn nngtnnnntt nnaaattatg tannatatnt ctt 533

<210> 1657  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1657

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 aaggggggtgg ctgggggtgct catggtggag atgngggggtc aggnaaagac nnnngggggg 120  
 gttgacagaa ntgggtgccta tattgtaagg cangctgcaa agantgncgt ggcaaatggc 180  
 cttgctagaa ngtgcattgt ncaagtttcc tatgcnattn gtgtncctaa nccntnntaa 240  
 atgttnttng atncnnattg aanttlnaan tanttcattn ataangataa ttanataat 300  
 taanaatnga ngaa 314

<210> 1658  
 <211> 557  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1658

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 ttcgggncga cccacgcgnc cggcnngccn canngaacca ccacacctct tntcgttttt 120  
 gctacccent tctgcnctac ggggaccggg naagtthtaaa nngggtaang atggcagaga 180  
 caatnnnatt taccnnagag tcggtgaacg agggacaccc tgacaagcnn tgcgacccaaa 240  
 tncccnatgc tgctctcnac gcttgccctng agcaggaccc anacagcaaa gngtgncctgc 300  
 tgaaacatgc accaaaacca actnggncat gggcttgga gaaatnacao ccaaggncac 360  
 ggcnagctac gacaagatag nncagagaca cctgcangaa cntnggacnt cgnntnaaat 420  
 gaagtngnga nttggatgcc nnaacttgc caagntccnn ngntaaccat tgtacaatna 480  
 tgatccccgg ngagttggtc naggcntaca agggncacn ntntcnanaa ancnggantt 540  
 attngnnnt tggtgnc 557

<210> 1659  
 <211> 285  
 <212> nucleic acid  
 <213> Glycine max

<400> 1659

cgctgcangc acgcgtacgt aagctcggaa ttccggctcga gnnactttcc tcttcacctc 60  
agaatctgta aacgaaggcc atcccgacaa gctgtgtgac caggtttcag atgccatcct 120  
agatgcatgc ttggagcaag acccagaaaag caaggttgct tgcgagacct gtacaaaaac 180  
taacatgggtt atgggtctttg gtgagattac aaccaaggcc agcgtgaact acgagaaaat 240  
agttcgagac acttgcaaag gcattggggtt tgtgtcacca gatgt 285

<210> 1660

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 1660

cacgcgtacg taagctcgga attcggctcg agcttcctct tcgcacaaaag cagcaagcat 60  
ccttgagatg gaaactttcc tcttcacctc agaatctgta aacgaaggcc atcccgacaa 120  
gctgtgtgac caggtttcag atgccatcct agatgcatgc ttggagcaag acccagaaaag 180  
caaggttgct tgcgagacct gtacaaaaac taacatgggtt atgggtctttg gtgagattac 240  
aaccaaggcc agcgtgaact acgagaaaat agttcgagac acttgcaaag gcattggggtt 300  
tgtg 304

<210> 1661

<211> 283

<212> nucleic acid

<213> Glycine max

<400> 1661

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ntccatcttc ttctttctctt 60  
cctcttcgca caaagcagca agnatecttg agatggaaac ttctctnctt cacctcagaa 120  
tctgtaaacg aaggccatcc cgacaagctg tgtgaccagg tttcagatgc catcctagat 180  
gcatgcttgg agcaagaccc agaaagcaag gttgcttgcg agacctgtac aaaaactaac 240  
atggttatgg tctttgggtga gattacaacc aaggccagcg tga 283

<210> 1662

<211> 447

<212> nucleic acid

<213> Glycine max

<400> 1662

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ccctcagcca tgtccttgca accaaacttg gtgctgcct cacagagggt aggaagaatg 120  
gcacctgtgc ttggttgagg ccagatggta agacacaagt aaccgtcgag tactacaatg 180  
acaatggtgc catggttcca gttcgtgtcc acactgtcct aatttccacc caacatgatg 240  
agactgtgag caatgatcaa attgctgagg accttaaaga gcatgttatc aagcctgtca 300  
ttcctgagaa gtaccttgat gagaagacca tcttccacct taacccttct ggccgttttg 360  
tcattgggtg ccctcatggt gatgctggtc tcaactggaa gaaagatnat cattgatacc 420  
taaggtggct ggggtgctca aggtgga 447

<210> 1663

<211> 475

<212> nucleic acid

<213> Glycine max

<400> 1663

ccacgcgtcc gcacaaagcg ggttactgtc tgttcaagct accatctctc tctctcttcc 60  
ttagngcctc cttgccagaa gttaaaatgg cccaagaaac tttcctattc acatctgaat 120  
cagtgaacga ggggcacctt gacaagctct gtgaccagat ctccgatgct gtgctcgatg 180  
catgcttgga gcaggacctt gacagcaagg ttgcctgtga aacctgcacc aagaccaaca 240  
tggtgatggt tttcggagag atcacaacca aggccaacgt ggactatgag aagattgtgc 300  
gtgacacatg caggaacatt ggttttgtct ctgatgatgt tggctttgat gctgacaact 360  
gcaaggctct cgtcaacatt gagcaacaga gtccatgatat tgctcaagggt gtgcacggnc 420  
acctnacaaa gaggcctgag gagattggtg ctggtgacca aggtcatatg ttccg 475

<210> 1664

<211> 520

<212> nucleic acid

<213> Glycine max

<400> 1664

gnngnnnnnn aggaggtntt gntntggaan cnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60

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nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn gagaagacga cagaaggggt 120
ctcaaagat gtgggactgg atgccgacaa ctgcaagggtc ctgctcaaca ttgagcagca 180
gagccctgat attgctcagg gtgtacacgg ccaccttacc aaaaaacctg aagaaattgg 240
tgctggtgac caggggtcaca tgtttggtta tgccactgat gaaacctctg aattgatgcc 300
attgagccat gttcttgcaa caaaactcgg tgctcgtctc accgagggtc gcaagaacgg 360
tacctgccct tggctgaggg ctgatgggaa gacccaagtg accgttgagt attacaatga 420
caatggtgcc aggggttcta ttctgtgaca caccgtgcta atctccacc aacacgacga 480
gactgtcacc aatgaccaa ttgntgntta acctnaaaaa 520

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<210> 1665
<211> 494
<212> nucleic acid
<213> Glycine max
<400> 1665

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aaactcgaa ttcggtcga gaacagcaca aagcgggtta ctgtctgttc aagctaccat 120
ctctctctct ctttcttagt gcctccttgc cagaagttaa aatggcccaa gaaactttcc 180
tattcacatc tgaatcagt aacgaggggc acctgacaa gctctgtgac cagatctccg 240
atgtgtgtgt cgatgcattg ttggagcagg acctgacag caagggttgc tgtgaaacct 300
gcaccaagac caacatggtg atgggttttc gagagatnac aaccaaggcc aacgtggact 360
atgagaagat tgtgcgtgac acatgcagga acattgggtt tgtctctgat gatgttggtc 420
ttgatgctga caactgcaag gtctcgtgca acattgagca acagagtcct gatattgctc 480
aaggtgtgca cngg 494

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<210> 1666
<211> 502
<212> nucleic acid
<213> Glycine max
<400> 1666

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gnagtgtttg ntntgggggg ggggagnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnaaagcgg gttactgtct 120

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gttcaagcta ccattctctct ctctctttct tagtgcctcc ttgccagaag ttaaaatggc 180  
ccaagaaact ttcctattca catctgaatc agtgaacgag gggcaccctg acaagctctg 240  
tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccctg acagcaaggt 300  
tgctgtgtaa acctgcacca agaccaacat ggtgatgggt ttccggagaga tcacaaccaa 360  
ggccaacgtg gactatgaga agattgtgcg tgacacatgc aggaacattg ggttttgtct 420  
ctgatgatgt tgggtcttgat gctgacaact gcaagggtccc tcgtcaacat tgagcaacag 480  
agtcttgata ttgctcaagg tg 502

<210> 1667  
<211> 372  
<212> nucleic acid  
<213> Glycine max  
<400> 1667

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acaagctctg tgaccagatn tccgatgctg tgctcgatgc atgcttggag caggacctga 120  
cagcaagggt gctgtgaaa cctgnaccaa gaccaacatg gtgatgggtt tcggagagat 180  
cacaaccaag gccaacgtgg actatgagaa gattgtgctg gacacatgca ggaacattgg 240  
ttttgtctct gatgatgttg gtcttgatgc tgacaactgc aagtcctcgt caacattgag 300  
caacagagtc ctgatattgc tcaagggtgtg cacggccact cacaaagagg cctgaggaga 360  
ttggtgtggt na 372

<210> 1668  
<211> 487  
<212> nucleic acid  
<213> Glycine max  
<400> 1668

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ttcggctcga ggtcgggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat 120  
gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc 180  
acaaaaacca acttgggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac 240  
gagaagatag tgcgtgacac ctgcaggaac atcggttcg tctcaaatga tgtgggactg 300

gatgccgaca actgcaaggt cctccgtcaa cattgagcag cagagccctg atattgctca 360  
 aggtgtacac gggcaactta ccaaaaaaacc tgaagaaatt ngtgctggtg accaaggtca 420  
 cattttgggt aatnccaactg gntgnaaacc ccngnattta tgncccattg accnagttct 480  
 tnncaaa 487

<210> 1669  
 <211> 419  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1669

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 tctctctctc tttcttagtg cctccttgcc agaagttaaa atggcccaag aaactttcct 120  
 attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc agatctccga 180  
 tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggttgct gtgaaacctg 240  
 caccaagacc aacatggtga tggttttcgg agagatcaca accaaggcca acgtggacta 300  
 tgagaagatt gtgcgtgaca catgcaggaa cattggtttt gtctctgatg atgttgggtc 360  
 ttgatgctga caactgcaag gtctctngtca acattgagca acagaatcct gatattgct 419

<210> 1670  
 <211> 447  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1670

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 ggtcaccocg acaagctgtg cgaccagatc tctgatgcag ngctcgatgc gtgccttgaa 180  
 caggaccctg acagcaaggt tgcctgtgag acatgcacca agaccaacat ggtcatggtc 240  
 tttggagaga tcacaaccaa ggccaacgta gactatgaga agattgtccg tgacacatgc 300  
 cgngaaattg gattcatctc tgatgatgtt ggtcttgatg ctgacaaatg caaggngttg 360  
 gtcaacattg aacancaaan ccctgatatc nccaggngn gcacggnac ttcaccaacc 420  
 cccaaaagaa ggttnggctn ggncca 447

<210> 1671  
 <211> 517  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1671

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 atgcacgcgt acgtaagctc ggaattcggc tcgagcacca caccactctc tctgctcttc 120  
 ttctaccttt naagnnttta aagtattaag atggcagaga cattcctatt tacctcagag 180  
 tcagtgaacg agggacaccc tgacaagctc tgcgacaaaa tctccgatgc tgtcctcgac 240  
 gcttgcccttg aacaggaccc agacagcaag gttgcctgcg aaacatgcac caagaccaac 300  
 ttggatcatgg tcttcggaga gatcaccacc aaggccaacg ttgactaccg aagaagatcc 360  
 gtgcgtgaca cctgcaggaa catcggcttc gtctcaaacg atgtgggact tgatgctgac 420  
 aactgcaagg tccttgtaaa cattgagcag cagagccctg atattgccca ggggtgtgcac 480  
 ggncacctta ccaaaagacc cgaggaaatc ggtgctg 517

<210> 1672  
 <211> 492  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1672

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 tgcaagaag acgacagaag ggggcaccgc ttgagcagac ttaacaacag cacaagcgg 120  
 gttactgtct gttcaagcta ccattctctc ctctctttct tagtgccctc ttgccagaag 180  
 ttaaaatggc ccaagaaact ttcctattca catctgaatc agtgaacgag gggcaccctg 240  
 acaagctctg tgaccagatc tccgatgctg tgctcgggtc atgcttgag caggaccctg 300  
 acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttcggagaga 360  
 tcacaaccaa ggccaacgtg gactatgaga agattgtgcg tgacacatgc aggaacattg 420  
 gttttgtctc tgatgatggt ggtcttgatg ctgacaactg caaaggctct cgtcaacatt 480  
 gagcaacaaa at 492

<210> 1673



<211> 503  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1673

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 cnaagaccaa agccccactc aaacaacaca ccaatctctc tgctcctcct cnaactttca 120  
 agttttttaa gtnttaaaga tggcagagac attcctaatt nacctcagag tcagtgaacg 180  
 agggacaccc tgacaagctc tgcgaccaaa tctccgatgc tgtcctcnac gcttgccttg 240  
 aacaggaccc agacagcaan gttgcctgcg aaacatgcac caagaccaac ttgggtcatgg 300  
 tcttcggaga gatnaccanc aaggccaacg ttgactacga gaagatngtg cgtgacacct 360  
 gcangaacat cggtttcgtc tcaaacgatg tgggacttga tgctgacaac tgcaagggtcc 420  
 ttgtaaacat tgagcaacaa aaccctgata ttgcccaagg tgtncacggc caacttacca 480  
 aaaganccga aggaaatcng tgc 503

<210> 1674  
 <211> 508  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1674

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 aggaacattg gttttgtctc tgatgatgtt ggtcttgatg ctgacaactg caaggctctc 180  
 gtcaacattg agcaacagag tcttgatatt gctcaagggtg tgcacggcca cctcaciaag 240  
 aggcctgagg agattggtgc tggtgaccaa ggtcatatgt tcggctatgc cactgacgag 300  
 actcccgagc tcatgccctt gagccatgtc cttgccacga agctcgggtgc caagctcanc 360  
 gacgggtccg aaaaacngga aatgcccttg ggctgaaaac ctgatggcaa nnaccaagtc 420  
 actgttggnn tactacaatt gacaaggggt ccatgggtcc aatccgcgtc aaaactgttg 480  
 ctcatntcca anacagcaat gatngaga 508

<210> 1675  
 <211> 334  
 <212> nucleic acid



acacgtgant aagcctgtga ttccctgagaa gtacctt

337

<210> 1678  
<211> 448  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1678

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gcaagtgtct tatgccattg gtgtgcctga gcctttgtct gtgtttgttg acacctatgg 120  
cactgggaag atccatgata aggagattct caacattgtg aaggaaaact ttgatttcag 180  
gcctggtatg atctccatca accttgatct caagaggggt ggaaataaca ggtttttgaa 240  
gactgctgcc tatggacact ttggaagaga agaccctgac ttcacatggg aagtggtaa 300  
accctcaag tgggagaagg cctaagtaat tcattccact gctctatgct ggaagttttt 360  
tgagcgttgc cttataata tgtctaata ccataacttt ccacgtctct tactctgtgt 420  
gtttctctcc tnttctcta ttttgga 448

Sequence 1

<210> 1679  
<211> 336  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1679

tgcangcgta cgtaagctcg gaattcggct cgagncgtct aagcagcatt gtggcaagtg 60  
gacttgccag aaggtgcatt gtgcaagtgt cttatgcat tgggtgcct gagcctttgt 120  
ctgtgtttgt tgacacctat ggcactggga agatccatga taaggagatt ctcaacattg 180  
tgaaggaaaa ctttgatttc aggctggta tgatctccat caaccttgat ctcaagaggg 240  
gtggaaataa cagggttttg aagactgctg cctatggaca ctttggaaga gaagacctg 300  
acttcacatg ggaagtggtc aaacctca antggg 336

<210> 1680  
<211> 493  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1680

gnnnnnnnga atgattngnt tontgocnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60  
 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn cccactcaac caccacacca ctctctctgc 120  
 tcttcttcta cctttcangt ttttacagta ttaagatggc agagacattc ctatttacct 180  
 cagagtcagt gaacgagggga caccctgaca agctctgcga ccaaactctcc gatgctgtcc 240  
 tcgacgcttg ccttgaacag gaccagaca gcaaggttgc ctgcgaaaca tgcaccaaga 300  
 ccaacttggt catggtcttc ggagagatca ccaccaaggc caacgttgac tacgagaaga 360  
 tcgtgcgtga cactgcagg aacatcggtc tcgtctcaaa cgatgtggga cttgatgctg 420  
 acaactgcaa ggtccttggt aacattgagc aacaaaacc tgataattgc caagggtttg 480  
 cacggccacc tta 493

Sequence:

<210> 1681  
 <211> 340  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1681

agtcgatgca cgcgtacgta agctcggaat tcngetcgag ntgccgcgaa attggattca 60  
 tctctgatga tgttggtctt gatgctgaca aatgcaaggt gttggtcaac attgagcaac 120  
 agagcccgga tatcgcccag ggtgtgcacg gccacttcac caagcgccca gaggagggtg 180  
 gtgctggtga ccagggtcac atgtttgggt atgccaccga tgaaaccccc gagtacatgc 240  
 cctcagcca tgtccttgca accaaacttg gtgctcgctt cacagagggt aggaagaatg 300  
 gcacctgtgc ttggttgagg ccagatggta agacacaagt 340

<210> 1682  
 <211> 317  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1682

cgt nagctcg gaattcggct cgagctgtga ccagatctcc gatgctgtgc togatgcatg 60  
 cttggagcag gaccctgaca gcaaaggttg cctgtgaaac ctgcaccaag accaactg 120  
 tgatggtttt cggagagatc acaaccaagg ccaacgtgga ctatgagaag attgtgcgtg 180  
 acacatgcag gaacattggt tttgtctctg atgaanttgg tcttgatgct gacaactgca 240

aggtcctcgt caacattgag caacagagtc ctgatattgc tcaaggtgtg cacggccacc 300  
tcacaaagag gcctgag 317

<210> 1683  
<211> 406  
<212> nucleic acid  
<213> Glycine max  
<400> 1683

gagaagacga cagaaggggg cagcgcttga gaccaagccc cactcaacca ccacaccact 60  
ctctctgctc ttcttctacc tttcaagttt ttaaagtatt aagatggcag agacattcct 120  
atttacctca gagtcagtga acgagggaca ccctgacaag ctctgcgacc aaatctccga 180  
tgctgtcctc gacgcttgcc ttgaacagga cccagacagc aaggttgcct gcgaaacatg 240  
caccaagacc aacttggtca tgggtcttcgg agagatcacc accaaggcca acgttgacta 300  
cgagaagatc gtgcgtgaca cctgcaggaa catcggttc gtctcaaacg atgtgggact 360  
tgatgctgac aactgcaagg tccttggttaa cattgagcaa caaaag 406

<210> 1684  
<211> 489  
<212> nucleic acid  
<213> Glycine max  
<400> 1684

actccaccgc gncggtaccg ttntaagncc ccgggccgac aaacgcgtca gtccgggctgc 60  
gagaagacga cagaaggggc accgcttgag cagacttaac aacagcacia agcgggttac 120  
tgtctgttca agctaccatc tctctctctc tttcttagtg ccttcttgcc agaagttaaa 180  
atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 240  
ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcanga ccctgacagc 300  
aaggttgcct gtgaaacctg caccaagacc aacatgggtga tggttttcgg agagatnaca 360  
accaagggca acgtggacta tgagaagatt gtgctgaca catgcaagaa cattgggtttt 420  
gtctctgatg aagttgggtc tgatgctgac aactgcaang gtctcgttca acattgcagc 480  
cancaagat 489

<210> 1685

<211> 506  
 <212> nucleic acid  
 <213> Glycine max

<400> 1685

gtggnnnnngg gnnaactttn aacngccang ggccggtana gaattaacgg ctcganccac 60  
 gcgtcaagta cggtgcgcnag aagacgacag aaggggatga aaccctgag tacatgcccc 120  
 tcagccatgt ccttgcaacc aaactcgggtg ctgcctcac cgaggtagg aaaaatggta 180  
 cctgtgcttg gctgaggcca gatggcaaga cacaagtaac tgttgagtac tacaatgaca 240  
 atggtgccat ggttccagtt cgtgtccaca ctgtcctaatt ttccacccaa catgttgaga 300  
 ctgtgagcaa tgaccaaatt gctgctgacc ttaaagaaca tgttatcaag cctgtcattc 360  
 ctgagaagtn cctggatgag aagaccatct tccaacctta aaccttctgg gcgtttttgn 420  
 cnnttggtgg gcccnnangg tganncccgg gcccanatgg gaaannaaag atttccccnt 480  
 ggaaaccocan aggttggnngn gggntc 506

<210> 1686  
 <211> 427  
 <212> nucleic acid  
 <213> Glycine max

<400> 1686

gaggccagge aagccccact caaccaccac acctctctc gggtcacgcc taccctttct 60  
 gctctttcttc nacctttcaa gttttaaaag tataaagatg gcanagacat toctattttac 120  
 ctgagagtgc gtgaacgagg gacacctga caagctctgc gaacaaatct ccgatgctgt 180  
 cctcgacgct tgctcgagc aggaccana cagcaaagtt gcctgcgaaa catgcaccaa 240  
 aaccaacttg gtcatggtct tcggagaaat cagaccaag gccaacgttg actacgagaa 300  
 gatagtgcgt gacacctgca ggaacatcgg cttcgtctca aatgatgtgg gactgggatg 360  
 ccgacaactg caaggtctct gtcaacattg agcancagan ccctgatatt gccanggtg 420  
 tacacgg 427

<210> 1687  
 <211> 504  
 <212> nucleic acid  
 <213> Glycine max

<400> 1687

ggnaactctt cgcggccaaa ctcttacann nocaggtagn gntanangaa ttcccggctc 60  
gacccacgcg tnacgtacgg ctgcgagaag acgacagaag ggggcagcgc ttgagaccaa 120  
gccccactca accaccacac cactctctct gctcttcttc tacctttcaa gtttttaaag 180  
tattaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 240  
caagctctgc gaccaaactct cccatgctgt cctcgacgct tgccttgaac aggaccaga 300  
cagcaagggt gcctgcgaaa catgcaccaa gaccaacttg gtcattgtct tcggagagat 360  
caccaccaag gccaacgttg actacganga gatcgtgcgt gacacctgca ggaacatcgg 420  
cttcgtctca aacgatgtgg gacttgatgc tgacaactgc aaggctccttg taaacattga 480  
agcagcagag ccctgatatt gccc 504

<210> 1688

<211> 323

<212> nucleic acid

<213> Glycine max

<400> 1688

ncgnangcac gcgtacgtna gctcgggaatt cggctcgagn ctgaatcngt gaacgagggg 60  
caccctganc aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttgagaca 120  
ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaactatgg tgatgggttt 180  
cggagagatc acaaccaagg ccaacgtgga ctatgagaag attgtgcgtg acacatgcag 240  
gaacattggg tttgtctctg atgatgttgg tcttgatgct gacaactgca aggtcctont 300  
caacattgag caacagagtc ctg 323

<210> 1689

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 1689

tcggctcgag ngaccatctt ccaccttaac ccttctggcc gttttgtcat tgggtggcct 60  
catgggtgatg ctgggtctcac tggaagaaa atcatcattg atacctatgg tggctggggg 120  
gctcatgggt gaggtgcctt ttcagggaag gaccctacca aggttgacag aagtgggtgcc 180

tatattgtaa ggcaggtgc aaagagtgtc gtggcaaata gccttgctag aaggtgcatt 240  
 gtgcaagttt cctatgccat tgggtgtccct gagcccttgt cagtgtttgt ggacac 296

<210> 1690  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1690

gcacgcgtac gtaagctcgg aattcggtc gagtatgaga agattgtccg tgacacatgc 60  
 cgcgaaattg gattcatctc tgatgatgtt ggtcttgatg ctgacaaatg caaggtgttg 120  
 gtcaacattg agcagcagag cctgatata gcccaggggtg tgcacgggtca cttaccaag 180  
 cgcccagagg aggttggtgc tgggtgaccag ggtcacatgt ttggctatgc cactgatgaa 240  
 acccctgagt acatgccct cagccatgtc cttgcaacca aactcgggtgc tcgcctcacc 300  
 gag 303

<210> 1691  
 <211> 336  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1691

gncnatccaa agcgtacgta agctcggaat tcggctcgag ggaacatcgg cttcgtctca 60  
 aatgatgtgg gactggatgc cgacaactgc aaggtcctcg tcaacattga gcagcagagc 120  
 cctgncattg ctcaggggtg acacggccac cttacaaaaa aacctgaaga aattgggtgct 180  
 ggtgaccagg gtcacatgtt tggctatgcc actgatgaaa ccctgaatt gatgccattg 240  
 agccatgttc ntgcaacaaa actcgggtgct cgtctcaccg aggttcgcaa gaacggtacc 300  
 tgcccttggc tgaggcctga tgggaagacc caagtg 336

<210> 1692  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1692

tcgcatgcnc gcgtacgtna gctcggaatt cggtcggagg ttaaaatggc ccaaganact 60



ttcctattca catctgaatc agtgaacgag gggcacccctg acaagctctg tgaccagatc 120  
 tccgatgctg tgctcgatgc atgcttggag caggaccctg acagcaagggt tgccctgtgaa 180  
 acctgcacca agaccaacat ggtgatgggtt ttccggagaga tcacaaccaa ggccaacgtg 240  
 gactatgaga agattgtgcg tgacacatgc aggaacattg gttttgtctc tgatgatggt 300  
 ggtcttgatg ctga 314

<210> 1693  
 <211> 321  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1693

gtcgcangca cgcgtacgta agctcggaat tcggctcgag tcattatcga tacttatgga 60  
 ggatgggggtg ctcatgggtg tgggtgctttc tccgggaagg accctaccaa ggttgatagg 120  
 agtgggtgctt acattgtgag acaggctgct aagagcattg tggcaagtgg acttgccaga 180  
 aggtgcattg tgcaagtgtc ttatgccatt ggtgtgcctg agcctttgtc tgtgtttggt 240  
 gacacctatg gcaactgggaa gatccatgat aaggagattc tcaacattgt gaaggaaaac 300  
 tttgatttca ggcttggat a 321

<210> 1694  
 <211> 514  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1694

gngnnnnagt ggntngtgna gtnnactnag nnaaatTTTtg naccggtcog gaattcccg 60  
 gtcgaccac gcgtccgtac gagaagacga cagaaggggg cagcgcttga tttgaggcca 120  
 ggcaagcccc actcaaccac cacacctctc ctcgttcag ctaccccttt ctgctcttct 180  
 tctacctttc aagtttttaa agtataaaga tggcagagac attcctatit acctcagagt 240  
 cgggtgaacga gggacaccct gacaagctct gcgaccaaT ctccgatgct gtcctcgacg 300  
 cttgcctcga gcaggaccca gacagcaaag ttgcctgcga aacatgcacc aaaaccaact 360  
 tggatcatggt cttcggagaa atcacgacca aggccaacgt tgactacgag aagatagtgc 420  
 gtgacacctg caggaacatc ggcttcgtct caaatgatgt gggactggat gccgacaact 480

514

Parameter	Value	Unit	Reference
Temperature	25.0	°C	[1]
Pressure	1.0	atm	[1]
Concentration	0.1	mol/L	[1]
Time	10.0	min	[1]
Wavelength	400.0	nm	[1]
Scan rate	1.0	nm/min	[1]
Slit width	1.0	nm	[1]
Detector	Si	nm	[1]
Source	Deuterium	nm	[1]
Monochromator	Grating	nm	[1]
Filter	None	nm	[1]
Cell	Quartz	nm	[1]
Path length	1.0	cm	[1]
Sample	Water	nm	[1]
Reference	None	nm	[1]
Wavenumber	2500.0	cm <sup>-1</sup>	[1]
Resolution	4.0	cm <sup>-1</sup>	[1]
Scan rate	1.0	cm <sup>-1</sup> /min	[1]
Slit width	1.0	cm <sup>-1</sup>	[1]
Detector	Si	cm <sup>-1</sup>	[1]
Source	Deuterium	cm <sup>-1</sup>	[1]
Monochromator	Grating	cm <sup>-1</sup>	[1]
Filter	None	cm <sup>-1</sup>	[1]
Cell	Quartz	cm <sup>-1</sup>	[1]
Path length	1.0	cm	[1]
Sample	Water	cm <sup>-1</sup>	[1]
Reference	None	cm <sup>-1</sup>	[1]

Parameter	Value	Unit	Reference
Temperature	25.0	°C	[1]
Pressure	1.0	atm	[1]
Concentration	0.1	mol/L	[1]
Time	10.0	min	[1]
Wavelength	400.0	nm	[1]
Scan rate	1.0	nm/min	[1]
Slit width	1.0	nm	[1]
Detector	Si	nm	[1]
Source	Deuterium	nm	[1]
Monochromator	Grating	nm	[1]
Filter	None	nm	[1]
Cell	Quartz	nm	[1]
Path length	1.0	cm	[1]
Sample	Water	nm	[1]
Reference	None	nm	[1]
Wavenumber	2500.0	cm <sup>-1</sup>	[1]
Resolution	4.0	cm <sup>-1</sup>	[1]
Scan rate	1.0	cm <sup>-1</sup> /min	[1]
Slit width	1.0	cm <sup>-1</sup>	[1]
Detector	Si	cm <sup>-1</sup>	[1]
Source	Deuterium	cm <sup>-1</sup>	[1]
Monochromator	Grating	cm <sup>-1</sup>	[1]
Filter	None	cm <sup>-1</sup>	[1]
Cell	Quartz	cm <sup>-1</sup>	[1]
Path length	1.0	cm	[1]
Sample	Water	cm <sup>-1</sup>	[1]
Reference	None	cm <sup>-1</sup>	[1]

Year	Age	Sex	Occupation	Education	Marital Status	Religion	Political Party	Income	Health	Smoking	Alcohol	Exercise	Stress	Family Size	Community	Environment	Genetics	Other
1990	25	M	Student	High School	Single	Catholic	Democrat	\$10,000	Good	Yes	No	None	Low	2	Urban	Good	None	
1991	26	F	Teacher	College	Married	Protestant	Republican	\$15,000	Good	No	Occasional	Regular	Low	3	Suburban	Good	None	
1992	27	M	Engineer	College	Single	Jewish	Democrat	\$20,000	Good	No	No	Regular	Low	1	Urban	Good	None	
1993	28	F	Nurse	College	Married	Catholic	Democrat	\$18,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
1994	29	M	Doctor	College	Single	Protestant	Republican	\$25,000	Good	No	No	Regular	Low	1	Urban	Good	None	
1995	30	F	Lawyer	College	Married	Jewish	Democrat	\$30,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
1996	31	M	Manager	College	Single	Catholic	Democrat	\$22,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
1997	32	F	Accountant	College	Married	Protestant	Republican	\$19,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
1998	33	M	Scientist	College	Single	Jewish	Democrat	\$28,000	Good	No	No	Regular	Low	1	Urban	Good	None	
1999	34	F	Writer	College	Married	Catholic	Democrat	\$21,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2000	35	M	Artist	College	Single	Protestant	Republican	\$17,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2001	36	F	Consultant	College	Married	Jewish	Democrat	\$24,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2002	37	M	Analyst	College	Single	Catholic	Democrat	\$23,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2003	38	F	Designer	College	Married	Protestant	Republican	\$20,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2004	39	M	Developer	College	Single	Jewish	Democrat	\$26,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2005	40	F	Executive	College	Married	Catholic	Democrat	\$29,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2006	41	M	Researcher	College	Single	Protestant	Republican	\$27,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2007	42	F	Manager	College	Married	Jewish	Democrat	\$25,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2008	43	M	Analyst	College	Single	Catholic	Democrat	\$24,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2009	44	F	Designer	College	Married	Protestant	Republican	\$22,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2010	45	M	Developer	College	Single	Jewish	Democrat	\$28,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2011	46	F	Executive	College	Married	Catholic	Democrat	\$30,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2012	47	M	Researcher	College	Single	Protestant	Republican	\$29,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2013	48	F	Manager	College	Married	Jewish	Democrat	\$27,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2014	49	M	Analyst	College	Single	Catholic	Democrat	\$26,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2015	50	F	Designer	College	Married	Protestant	Republican	\$25,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2016	51	M	Developer	College	Single	Jewish	Democrat	\$28,000	Good	No								

Year	Age	Sex	Occupation	Education	Marital Status	Religion	Political Party	Income	Health	Smoking	Alcohol	Exercise	Stress	Family Size	Community	Environment	Genetics	Other
1990	25	M	Student	High School	Single	Catholic	Democrat	\$10,000	Good	Yes	No	None	Low	2	Urban	Good	None	
1991	26	F	Teacher	College	Married	Protestant	Republican	\$15,000	Good	No	Occasional	Regular	Low	3	Suburban	Good	None	
1992	27	M	Engineer	College	Single	Jewish	Democrat	\$20,000	Good	No	No	Regular	Low	1	Urban	Good	None	
1993	28	F	Nurse	College	Married	Catholic	Democrat	\$18,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
1994	29	M	Doctor	College	Single	Protestant	Republican	\$25,000	Good	No	No	Regular	Low	1	Urban	Good	None	
1995	30	F	Lawyer	College	Married	Jewish	Democrat	\$30,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
1996	31	M	Manager	College	Single	Catholic	Democrat	\$22,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
1997	32	F	Accountant	College	Married	Protestant	Republican	\$19,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
1998	33	M	Scientist	College	Single	Jewish	Democrat	\$28,000	Good	No	No	Regular	Low	1	Urban	Good	None	
1999	34	F	Writer	College	Married	Catholic	Democrat	\$21,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2000	35	M	Artist	College	Single	Protestant	Republican	\$17,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2001	36	F	Consultant	College	Married	Jewish	Democrat	\$24,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2002	37	M	Analyst	College	Single	Catholic	Democrat	\$23,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2003	38	F	Designer	College	Married	Protestant	Republican	\$20,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2004	39	M	Developer	College	Single	Jewish	Democrat	\$26,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2005	40	F	Executive	College	Married	Catholic	Democrat	\$29,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2006	41	M	Researcher	College	Single	Protestant	Republican	\$27,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2007	42	F	Manager	College	Married	Jewish	Democrat	\$25,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2008	43	M	Analyst	College	Single	Catholic	Democrat	\$24,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2009	44	F	Designer	College	Married	Protestant	Republican	\$22,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2010	45	M	Developer	College	Single	Jewish	Democrat	\$28,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2011	46	F	Executive	College	Married	Catholic	Democrat	\$30,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2012	47	M	Researcher	College	Single	Protestant	Republican	\$29,000	Good	No	No	Regular	Low	1	Urban	Good	None	
2013	48	F	Manager	College	Married	Jewish	Democrat	\$27,000	Good	No	No	Regular	Low	2	Suburban	Good	None	
2014	49	M	Analyst	College	Single	Catholic	Democrat	\$26,000	Good	No	Occasional	Regular	Low	1	Urban	Good	None	
2015	50	F	Designer	College	Married	Protestant	Republican	\$25,000	Good	No	Occasional	Regular	Low	2	Suburban	Good	None	
2016	51	M	Developer	College	Single	Jewish	Democrat	\$28,000	Good	No								

aactcttacg tngccaggnn ccggtanaga attaccgggg ncgacccacg cgtcngccca 60  
cncgtccgcc cagcggtccg acggctgcga gaagacgaca gaaggggggc agcgcttgag 120  
accaagcccc actcaaccac cacaccactc tctctgctct tcttctacct ttcaagtttt 180  
taaagtatta agatggcaga gacattccta tttacctcag agtcagtga cagaggacac 240  
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgcc tgaacaggac 300  
ccagacagca aggttgccctg cgaaacatgc accaagacca acttggtcat ggtcttcgga 360  
gagatnacca ccaaggccaa cgttgactac gagaagatcg tgcgtgacac ctgcaggaac 420  
atcggttcg tctcaaacga tgtgggactg atgctgacaa ctgcaaagtc cttgttaaca 480  
atgaacanca aanccc 496

1698  
300  
nucleic acid  
Glycine max  
1698  
60  
120  
180  
240  
300

<210> 1698  
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<212> nucleic acid  
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<400> 1698  
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gcaagctgca aagagcattg ttgcaaattg acttgctagg agggcaattg tgcaagtttc 120  
ctatgccatt ggtgtgcctg agcccttgtc tgtgtttgtt gacacttatg gcactgggaa 180  
gatccctgac aaggaaatcc tcagcattgt gaaggagagt tttgaacttca ggccctggcat 240  
gatctccatc aaccttgatc tcaagagggg tggaaatggc aggttccttga agactgctgc 300

<210> 1699  
<211> 303  
<212> nucleic acid  
<213> Glycine max  
<400> 1699

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cacaaccaag gccaacgtag actatgagaa gattgtccgt gacacatgcc gcgaaattgg 120  
attcatctct gatgatgttg gtcttgatgc tgacaaatgc aagggtgttg toaacattga 180  
gcagcagagc cctgatatcg ccagggtgt gcacggtcac ttcaccaage gccagagga 240  
ggttggtgct ggtgaccagg gtcacatggt tggctatgcc actgatgaaa cccctgagta 300

cat

303

<210> 1700  
<211> 311  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1700

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tggtttttcgg agagatcaca accaaggcca acgtggacta tgagaagatt gtgcgtgaca 120  
catgcaggaa catttggtntt gtctctgatg atgttggtct tgatgctgac aactgcaagg 180  
tcctcgtaaa cattgagcna cagagtcctg atattgctca aggtgtgcac gnccacctca 240  
caaagaggcc tgaggagatt ggtgctggtg accaaggta tatgttcggc tatgccactg 300  
acgagactcc c 311

<210> 1701  
<211> 425  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1701

gagtaacaac agcacaaagc gggttactgt ctgttcaagc taccatctct ctctctcttt 60  
cttagtgcct ccttgccaga agttaaaatg gcccaagaaa ctttcctatt cacatctgaa 120  
tcagtgaacg agggggcacc tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat 180  
gcatgcttgg agcaggaccc tgacagcaag gttgcctgtg aaacctgcac caagaccaac 240  
atggtgatgg ttttcggaga gatcacaacc aaggccaacg tggactatga gaagattgtg 300  
cgtgacacat gcaggaacat tggttttgtc ccgatgatgt ttggtcctga tgctgacaac 360  
tgcaangtcc cccgtcaaca atgagcaaca nagtcctga aaattgcna angngttgna 420  
cgggc 425

<210> 1702  
<211> 321  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1702

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aaggtgcatt gtgcaagtnt cttatgccat tgggtgtgcc gagcctttgt ctgtctttnt 120  
tgacacctat ggcaccgggn agatccatga taaggagatt ctnacattg tgaaggagaa 180  
ctttgatttc aggcccggta tgatctccat caaccttgat ctcaagaggg gtgggnataa 240  
caggttcttg aagactgctg catatggncat cttcggcaga gaggaccctg acttcacatg 300  
ggaagtggtc nagccccctca a 321

<210> 1703  
<211> 311  
<212> nucleic acid  
<213> Glycine max

<400> 1703  
tcgcangcac gentacgtaa gctcgggaatt cggctcgagc ggctcgaggt agactatgag 60  
aagattgtcc gtgacacatg ccgcgaaatt ggattcatct ctgatgatgt tggctcttgat 120  
gctgacaaat gcaagggtgtt ggtcaacatt gagcagcaga gccctgatat cggccagggt 180  
gtgcacggtc acttcaccaa gcgcccagag gaggttgggt ctggtgacca gggtcacatg 240  
tttggtatg ccaactgatga aacccttgag tacatgcccc tcagccatgt ccttgcgccc 300  
aaactcgggtg n 311

<210> 1704  
<211> 473  
<212> nucleic acid  
<213> Glycine max

<400> 1704  
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gctgcgagaa gacgacagaa gggggcagcg cttganacca agccccactc aaccaccaca 120  
ccactctctc tgctcttctt ctaccttca agtttttaaa gtantaagat ggcagagaca 180  
ttcctattta cctcanagtc agtgaacgag ggacaccctg acaagctctg cgaccaaata 240  
tcgatgctg tctcgcagc ttgccttgaa caggaccag acagcaaggt tgctgcgaa 300  
acatgcacca ngaccanctt ggtcatgggt cttcggagag atcaccacca aggccaacgt 360  
tgactacgag aagatcgtgc gtgacacctg caggaacatc ggcttcgtct caaacgatgt 420

gggacttgat gctgacaact gcaaggtcct tgtaaacatt gagcagcaga gcc 473

<210> 1705  
 <211> 319  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1705

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 aatgggtgcc aatgttccagt tcgtgtccac actgtcctaa tttccacca acatgatgat 120  
 tctgtgagca atgaccaa atgctgtgac cttaaagagc atgttatcaa gcctgtcatt 180  
 cctgagaagt acctggatga gaagaccatc ttccaacctt aacccttctg gccgttttgt 240  
 cattggtggc cctcatgggtg atgctgggtct cactggaaga aagatcatca ttgataccta 300  
 tgggtgggtgg ggtgctcat 319

<210> 1706  
 <211> 507  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1706

gnnnnnnaan tctacgccgc cctcctaacg ngcacaanat tcncgggaac gacccaacgcg 60  
 nccgtacggc tgcaagaag acgacagaag ggggcagcgc ttgagancaa nccccactca 120  
 accaccacac cactctctct gctcttcttc nantttcaa gtttttaaag tattaagatg 180  
 gcagagacat tctatattac ctgagagtca gtgaacgagg gacaccctga caagctctgc 240  
 gacaaaatct ccgatgctgt cctcgacgct tgcctttaaa cangacccaa gacagcaaag 300  
 ttgcctgcca aacatgcacc aagaccaact tggatcatggt ctccggagag atnaccacca 360  
 agggcaacgt tgactacgag aagatcgtgc gtgacacctg caggaacatc ggcttcgtct 420  
 caaacgatgt gggacttgat gctgacaact gcaangtcct tgtaaacatt gagcaacaaa 480  
 accctganaa tncccaaggt ttcaccg 507

<210> 1707  
 <211> 351  
 <212> nucleic acid  
 <213> Glycine max

<400> 1707

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ctgcaaagag tgtcgtggca aatggccttg ctagaagggtg cattgtgcaa gtttcctatg 120  
ccattgggtgt ccctgagccc ttgtcagtgt ttgtggacac ttatggaact ggaagattc 180  
ctgacaagga gattctgcaa attgtgaagg agaatttcga cttcagacct ggaatgatca 240  
ccattaactt ggaccttaag aggggtgggc atagggtcct caagacagtg cttatggaca 300  
ctttggaagg gatgatgcag cttcactggg aagtgtgaac cactcaagtc a 351

<210> 1708

<211> 509

<212> nucleic acid

<213> Glycine max

<400> 1708

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gncgacncac gcgtccnccc acgcgtccgc ccacgcgtcc gccacgcgt ccgctgcgag 120  
aagacgacag aagggggcag cgcttgagac caagccccac tcaaccacca caccactctc 180  
tctgctcttc ttctaccttt caagttttta aagtattaag atggcagaga cattcctatt 240  
tacctcagag tcagtgaacg agggacaccc tgacaagctc tgcgacacaaa tctccgatgc 300  
tgtcctcgac gcttgcccttg aacaggaccc agacagcaag gttgcctgcg aaacatgcac 360  
caagaccaac ttggatcatgg tcttcggaga gatcaccacc aaggccaacg ttgactacga 420  
gaagatcgtg cgtgacacct gcaggaacat cggcttcgtc tcaaacgatg tgggacttga 480  
tgttgacaac tgcaagggtcc ttgtnaaca 509

<210> 1709

<211> 267

<212> nucleic acid

<213> Glycine max

<400> 1709

gagacaggct gctaagagca ttgtggcaag tggacttgcc agaagggtgca ttgtgcaagt 60  
gtcttatgcc attgggtgtgc ctgagccttt gtctgtgttt gttgacacct atggcactgg 120  
gaagatccat gataaggaga ttctcaacat tgtgaaggaa aactttgatt tcaggcctgg 180

tatgatctcc atcaaccttg atctcaagag ggggtggaaat aacaggtttt tgaagactgc 240  
 tgcctatgga cacttttgaa gagaaga 267

<210> 1710  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1710

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 cgcccagggt gtgcacggtc acttcaccaa gcgccagag gaggttggtg ctggtgaccn 120  
 cggtcacatg tttggctatg cactgatga aaccctgag tacatgcccc tcagccatgt 180  
 ccttgcaacc aaactcgggtg ctgcctcac cgaggtagg aaaaatggta cctgtgcttg 240  
 gctgaggcca gatggcaaga cacaagtaac tgttgagtac tacaatgaca atggtgccat 300  
 ggttccagtt cgtgtccaca 320

<210> 1711  
 <211> 330  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1711

nnnaaaannt gacntcgcan gcacggtac gtaagctcgg aattcggctc gagggactgg 60  
 atgccgacaa ctgcaaggtc ctgctcaaca ttgagcagca gagccctgat attgctcagg 120  
 gtgtacacgg ccaccttacc aaaaaacctg aagaaattgg tgctggtgac cagggtcaca 180  
 tgtttggcta tgccactgat gaaaccctg aattgatgcc attgagccat gttcttgcaa 240  
 caaaactcgg tgctcgtctc accgaggttc gcaagaacgg tacctgccct tggtgagge 300  
 ctgatgggaa gaccaagtg accgttgagt 330

<210> 1712  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1712

agtngcangc acgcgtacgt aagctcggaa ttcggctcga gtgcaagttt octatgccat 60



tgggtgtgcct gagcccttgt ctgtgtttgt tgacacttat ggcactggga agatccctga 120  
 caaggaaatc ctcagcattg tgaaggagag ttttgacttc aggcctggca tgatctccat 180  
 caaccttgat ctcaagaggg gtggaaatgg caggttcttg aagactgctg catatggaca 240  
 ctttggcaga gatgaccctg acttcacatg ggaagtgggtg aagccactca agggggagaa 300  
 ggtacctgct tac 313

<210> 1713  
 <211> 486  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1713

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 agaagacgac agaagggggc agcgcttgat ttgaggccag gcaagcccca ctcaaccacc 120  
 acacctctcc tcgttcaogc taccoccttc tgctcttctt ctacctttca agtttttaaaa 180  
 gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg 240  
 acaagctctg cgaccaaate tccgatgctg tctcgcagc ttgcctcgag caggaccacg 300  
 acagcaaagt tgcttgcgaa acatgcacca aaaccaactt ggtcatggtc ttcggagaaa 360  
 tcacgaccaa ggccaacgtt gactacgaga agatagtgcg tgacacctgc angaacatcg 420  
 gcttcgtctc aaatgatgtg ggactggatg ccgacaactg caaggtcctc gtcaacattg 480  
 agcaac 486

<210> 1714  
 <211> 474  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1714

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 gctgcgagaa gacgacagaa gggggcagcg cttgagacca agcccaactc aaccaccaca 120  
 ccactctctc tgctcttctt ctacctttca agtttttaaa gtattaagat ggcagagaca 180  
 ttctatttta cctcagagtc agtgaacgag ggacaccctg acaagctctg cgaccaaate 240  
 tccgatgctg tctcgcagc ttgccttgaa caggaccacg acagcaaggt tgcttgcgaa 300

acatgcacca agaccaactt ggtcatggtc ttcggagaga tcaccaccaa ggccaacgtt 360  
gactacgana agatcgtgcg tgacacctgc aggaacatcg gcttcgtctc aaacgatgtg 420  
ggacttgatg ctgacaactg caaaggctct ttgtaaacad tgagcaacan agnc 474

<210> 1715  
<211> 382  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1715

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tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180  
aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac 240  
agcaagggtg cctgtgaaac ctgcaccaag accaacadtg tgatgggtttt cggagagatc 300  
acaaccaagg ccaacgtgga ctatgagaag attgtgctg acacatgcag gaacatttgt 360  
tttgtctctg atgatgttgg tc 382

<210> 1716  
<211> 308  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1716

nntcgcangc acgcgtacgt aagctcggaa ttcggctcga gggcacccctg nacaagctct 60  
gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 120  
ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag atcacaacca 180  
aggccaacgt ggactatgag aagattgtgc gtgacacatg caggaacatt ggttttgtct 240  
ctgatgatgt tggctcttgat gctgacaact gcaaggctct cgtcaacatt gagcaacaga 300  
gtcctgat 308

<210> 1717  
<211> 312  
<212> nucleic acid  
<213> Glycine max

<400> 1717

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tcgatgcatg cttggagcag gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaaga 120  
ccaacatggt gatggttttc ggagagnnca caaccaaggc caacgtggac tatgagaaga 180  
ttgtgcgtga cacatgcagg aacattggtt ttgtcnctga tgatgttggc cttgatgctg 240  
acaactgcaa ggtcctcgtc aacattgagc aacagagtcc tgatattgct caagggtgtgc 300  
cgccacctc ac 312

<210> 1718

<211> 315

<212> nucleic acid

<213> Glycine max

<400> 1718

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catgtttggc tatgccactg atgaaacccc agaattcatg ccattgagtc atgttcttgc 120  
aaccaagctc ggtgctcgtc tcaccgaggt tcgcaagaac ggaacctgcc catggctgag 180  
gcctgatggg aagacccaag tgactgtgga gtattacaat gataatggtg ccaggggtcc 240  
agttcgtggt cacaccgtgc taatctccac ccagcatgat gagactgtca ccaacgacga 300  
aattgoggct gacct 315

<210> 1719

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 1719

ggaattcggc tcgagctggt gagtactaca atgacaatgg tgccatggtt ccagttcgtg 60  
tcacactgt cctaatttcc acacaacatn anaaggngt gagcaatgac caaattgctg 120  
ctgaccttaa agagcatggt atcaagcctg tnantcctga gaagtacctg gatgagaaga 180  
ccatctttna ccttaacctt tctggccgtt ttgtcattgg tggccctcat ggtgatgctg 240  
gtctcatgga agaaagatca tcattgatac ctatggtggg tggggtgctc atggtggagg 300  
tgcttttcag gg 312

<400>	1720
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[illegible]

<400>	1721
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[illegible]

<400>	1722
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gatgatgttg gtcttgatgc tgacaaatgc aaggtgttgg tcaacattga gcaacagagc 240  
 ccggatatcg cccaggggtgt gcacggccat tcaccaagcg cccagaggag gttggtgccg 300  
 gtgaccag 308

<210> 1723  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1723

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 atgcttggag caggaccctg acagcaaggt tgcctgtgaa acctgcacca agaccaacat 180  
 ggtgatggtt ttccgagaga tcacaaccaa ggccaacgtg gactatgaga agattgtgcg 240  
 tgacacatgc aggaacattg gtttgtctct gatgatgttg gtccatgatgc 290

<210> 1724  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1724

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 ccaacgttga ctacgagaag atcgtgcgtg acacctgcag gaacatcggc ttcgtctcaa 180  
 acgatgtggg acttgatgct gacaactgca aggtccttgt aaacattgag cagcagagcc 240  
 ctgatattgc ccaggggtgtg cacggccacc ttacaaaaag acccgaggaa atcgggtgctg 300  
 gagaccaggg tcacatgttt ggcta 325

<210> 1725  
 <211> 486  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1725

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cgacccacgc gtcngacggc tgcgagaaga cgacagaagg gggcagcgct tgagaccaag 120  
 ccccaactcaa ccaccacacc actctctctg ctcttcttct acctttcaag tttttaaaagt 180  
 attaagatgg cagagacatt cctatttacc tcagagtcag tgaacgaggg acaccctgac 240  
 aagctctgcg accaaatctc cgatgctgtc ctogacgctt gccttgaaca ggaccagac 300  
 agcaagggtg cctgcgaaac atgcaccaag accaacttgg tcatgggtctt cggagagatc 360  
 accaccaagg ccaacgttga ctacgagaag atcgtgcgtg acacctgcag gaacatcggc 420  
 ttcgtctcaa acgatgtggg acttgatgct gacaactgca aggtcccttg taaacattga 480  
 nncagc 486

<210> 1726  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1726

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 ctgaggagat tgggtgctggg gaccaaggtc atatgttcgg ctatgccact gacgagactc 120  
 ccgagctcat gcccttgagc catgtccttg ccacgaagct cgggtgccaag ctcaccgagg 180  
 ttcggaagaa cgggacatgc ccttgggtga gacctgatgg caagacccaa gtcaactgtt 240  
 agtactacaa tgacaagggg gccatgggtc caatccgcgt ccacactgtg ctcatctcca 300  
 cacagcat 308

<210> 1727  
 <211> 307  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1727

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 ctctgatgat gttgggtctt atgctgacaa ctgcaaggtc ctcgtcaaca ttgagcaaca 180  
 gagtcctgat attgctcaag gtgtgcacgg ccacctcaca aanaggcctg aggagattgg 240  
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cttgagc

307

<210> 1728  
<211> 313  
<212> nucleic acid  
<213> Glycine max

<400> 1728

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gagcctttgt ctgtgtttgt tgacaacctat ggcactgna agatccatga taaggagatt 180  
ctcaacattg tgaaggaaaa ctttgatttc aggcctggta tgatctccat caaccttgat 240  
ctcaagaggg gtggaaataa cagggtttttg aagactgctg cctatggaca ctttgggaaga 300  
gaagacctga ctt 313

<210> 1729  
<211> 320  
<212> nucleic acid  
<213> Glycine max

<400> 1729

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ttcatctctg atgatgttgg tcttgatgct gacaaatgca aggtgttggg caacattgag 180  
caacagagcc cggatatcgc ccagggtgtg cacggccact tgcaccaagc gccagagga 240  
ggttggtgct ggtgaccagg gtcacatgtt tgggtatgcc accgatgaaa ccccgagta 300  
catgcccctc agccatgtcc 320

<210> 1730  
<211> 361  
<212> nucleic acid  
<213> Glycine max

<400> 1730

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tgcacnacgc tgctttcttn ngacntgagg aatgggcaca agaaaccttt tctattcaca 120

tctgaatctg taaacgaggg ttcaccccga caagctgtgc gaccagatct ctgatgcagt 180  
gctcgatgcg tgccttgaac aggaccctga cagcaaggtt gcctgtgaga catgcaccaa 240  
gaccaacatg gtcatggtct ttgggagaga tcacaaccaa ggccaacgta gactatgaga 300  
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c 361

<210> 1731  
<211> 327  
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<213> Glycine max

<400> 1733

ngtcgcangc acgcgtacgt aagctcggaa ttcggctcga gattgatacc tatggtgggt 60  
ggggtgctca tgggtggaggt gccttttcag ggaaggaccc taaccaaggt tgacagaagt 120  
ggtgcctata tcgtgaggca ggctgcaaag agtggttggt caaatggcct tgccagaaggt 180  
tgcattgtcc aagtttctta tgccattggt gtccctgagc ccttgtcagt gtttgtggac 240  
acttatggaa ctgggaagat tcctgacaag gagattcttc aaattgtgaa ggagaatttc 300  
gatttcagac ctggaatgat c 321

<210> 1734

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 1734

gtcgcnngca cgcgtacgta agctcggaa ttcggctcga ggacgagacc ccagaattga 60  
tgccattgag tcatgttctt gcaactaaac tccgtgctcg tctcaccgag gttcgcaaga 120  
acggaacctg cccatggttg aggcctgatg ggaagaccca agtgactgtt gagtattaca 180  
atgacaacgg tgccatggtt ccagttcgtg tccacactgt gcttatctcc acccaacatg 240  
atgagactgt gaccaacgac gaaattgcag ctgacctcaa ggagcatgtg atcaagccgg 300  
tgatcccgga 310

<210> 1735

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 1735

ngtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gtggtcatgg tctttggaga 60  
gatcacaacc aaggccaacg tagactntga gaagattgtc cgtgacacat gccgcgaaat 120  
tggaattcatc tctgatgatg ttggtcttga tgctgacaaa tgcaagggtg ttgtcaacat 180  
tgagcagcag agccctgata tcgcccaggg tgtgcacggt cacttcacca agcgcccaga 240  
ggaggttggt gctggtgacc agggtcacat gtttggctat gccactga 288

<210> 1736  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1736  
  
 gtcgcangca cgcgtacgtg agctcgggnt tcggctcgag ntcagtgaac gaggggcacc 60  
 ctgacaagct ctgngaccag atctccgatg ctgtgctcga tgcattgctt gagcaggacc 120  
 ctgacagcaa ggttgccctgt naaacctgca ccaagaccaa catggtgatg gttttcggag 180  
 agatcacaac caaggccaac gtggactatg agaagattgt gcgtgacaca tgcaggaaca 240  
 ttggttttgt ctctgatgat gttggtcttg atgctgacaa ctgcaaggtc ctcnncaan 299

<210> 1737  
 <211> 328  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1737  
  
 ngtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggccgtgcat tcctgagaag 60  
 taccttgatg agaagaccat cttccacctt aacccttctg gccgttttgt cattggtggc 120  
 cctcatggtg atgctggtct cactggaaga aagatcatca ttgataccta tggaggctgg 180  
 ggtgctcatg gtggagggtgc cttttcaggg aaggacccta ccaagggtga cagaagtgg 240  
 gcctatattg taaggcaggc tgcaaagagt gtcgtggcaa atggccttgc tagaagggtgc 300  
 ttgtgcaagt ttccctatgc catggtgc 328

<210> 1738  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1738  
  
 tcncgcgtac gtnagctcgg aattcggctc gagaccaaga ccaacatggg gatgggtttc 60  
 ggagagatca caaccaaggc caacgtggac tatgagaaga ttgtgcgtga cacatgcagg 120  
 aacattggtt ttgtctctga tgatgttggt cttgatgctg acaactgcaa ggtcctcgtc 180  
 aacattgagc aacagagtcc tgatattgct caagggtgtgc acggccacct tcacaaagag 240

gcctgaggag attggtgctg gtgaccaagg tcatatgttc ggctatgccc actgacgaga 300  
ctcccagact cagcc 315

<210> 1739  
<211> 303  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1739

cccngcacgc gtacgtaagc tcggaattcg gctcgagcga tacttatgga ggatggggtg 60  
ctcatgggtg tggtgctttc tcogggaagg accctaccaa ggttgatagg agtgggtgctt 120  
acattgtgag acaggctgct aagagcattg tggcaagtgg acttgccaga aggtgcattg 180  
tgcaagtgtc ttatgccatt ggtgtgcttg agcctttgtc tgtgtttgtt gacacctatg 240  
gcactgggaa gatccatgat aaggagattc tcaacattgt gaaggaaaac tttgattcag 300  
gcc 303

<210> 1740  
<211> 299  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1740

tctntgnanc gtagtaagct cggaattcgg ctcgagctga tattgcccag ggtgtgcacg 60  
gccaccttac caaaagaccc gaggaatcg gtgctggaga ccagggtcac atgtttggct 120  
atgccacgga cgagacccca gaattgatgc cattgagtca tgttcttgca actaaactcg 180  
gtgctcgtct caccgagggt cgcaagaacg gaacctgccc atgggtgagg cctgatggga 240  
agacccaagt gactgttgag tattacaatg acaacggtgc catggttcca gttcgtgtc 299

<210> 1741  
<211> 263  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1741

cattgagcaa cagagtcctg atattgctca aggtgtgcac ggccacctca caaagaggcc 60  
tgaggagatt ggtgctggtg accaagggtca tatgttcggc tatgccactg acgagactcc 120

cgagctcatg cccttgagcc atgtccttgc cacgaagctc ggtgcccaagc tcaccgaggt 180  
 tcggaagaac gggacatgcc cttggctgag acctgatggc aagaccaag tcactgttga 240  
 gtactacaat gacaaggggtg cca 263

<210> 1742  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1742

gtngangcgt acgtaagctc ggaattcggc tcgaggcacg gccacctcac aaagaggcct 60  
 gaggagattg gtgctgggtga ccaaggtcat atgttcggct atgccactga cgagactccc 120  
 gagctcatgc ccttgagcca tgtccttgcc acgaagctcg gtgccaaagct caccgaggtt 180  
 cggaagaacg ggacatgccc ttggctgaga cctgatggca agaccaagt cactgttgag 240  
 tactacaatg acaaggggtgc catggttcca atccgcgtcc aactgtgtct catctccac 299

<210> 1743  
 <211> 254  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1743

ctcaccgagg ttcgcaagaa cggtagctgc ccttggttga ggctgatgg gaagacccaa 60  
 gtgaccgttg agtattacaa tgacaatggt gccagggttc ctattcgtgt acacaccgtg 120  
 ctaatctcca cccaacacga cgagactgtc accaatgacg aaattgctgc tgacctcaaa 180  
 gagcatgtga tcaagcctgt gatcccagag aagtaccttg atgagaagac cattttccac 240  
 ttgaaccctt cagg 254

<210> 1744  
 <211> 268  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1744

acagagtect gatattgctc aagggtgtgca cgccacctc acaaagaggc ctgaggagat 60  
 tgggtgctggg gaccaaggtc atatgttcgg ctatgccact gacgagactc ccgagctcat 120

gcccttgagc catgtccttg ccacgaagct cggtgccaag ctcaccgagg ttcggaagaa 180  
 cgggacatgc ccttggtgta gacctgatgg caagacccaa gtcactgttg agtactacaa 240  
 tgacaagggt gccatggttc caatccgc 268

<210> 1745  
 <211> 305  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1745

gcacgcgtac gtaagctcgg aattcggctc gagcacggac gagaccccag aattgatgcc 60  
 attgagtcac gttcttgcaa cttaaactcgg tgctcgtctc accgaggttc gcaagaacgg 120  
 aacctgcccc tgggttgaggc ctgatgggaa gacccaagtg actgttgagt attacaatga 180  
 caacgggtgcc atggtttccag ttcgtgncca cactgtgctt atctccaccc aacatgatga 240  
 gactgtgacc aacgacgaaa ttgcagctga cctcaaggag catgtgatca agccggtgat 300  
 cccgg 305

<210> 1746  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1746

antcgcangc acgcgtacgt aagctcggaa ttcggctcga ggtcctcgac gcttgccctg 60  
 aacaggaccc agacagcaag gttgcctgcg aaacatgcac caagaccaac ttggtcatgg 120  
 tcttcggaga gatcaccacc aaggccaacg ttgactacga gaagatcgtg cgtgacacct 180  
 gcaggaacat cggcttcgtc tcaaacgatg tgggacttga tgctgacaac tgcaagggtcc 240  
 ttgtaaacat tgagcagcag agccctgata ttgccagggt tgtgcacggc caccttacca 300  
 aaagacccga ggaaat 316

<210> 1747  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1747

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ctcaccggcc gcaagatcat 60  
catcgacacc tatggaggat ggggtgcaca tgggtggtggt gccttctctg ggaaggatcc 120  
taccaagggtt gataggagtgtgtgcctacat tgtgaggcaa gctgcaaaga gcattgttgc 180  
aatggactt gctaggaggg caattgtgca agtttctctat gccattggtg tgcctgagcc 240  
cttgtctgtg tttgttgaca cttatggcac tgggaagatc cctgacaagg aaatcctcag 300  
cattgt 306

<210> 1748  
<211> 269  
<212> nucleic acid  
<213> Glycine max

<400> 1748

gtgcgtgaca catgcaggaa cattgggtttt gtctctgatg atgttggtct tgatgctgac 60  
aactgcaagg tcctcgtcaa cattgagcaa cagagtcctg atattgctca aggtgtgcac 120  
ggccacctca caaagaggcc nnaggagatt ggtgctggtg accaagggtca tatgttcggc 180  
tatgccactg acgagactcc cgagctcatg cccttgagcc atgtccttgc cacgaagctc 240  
ggtgccaagc tcaccgaggt tcggaagaa 269

<210> 1749  
<211> 311  
<212> nucleic acid  
<213> Glycine max

<400> 1749

tcgcangcac ncgtacgtaa gctcggaatt cggctcgaga cagagtcctg atattgctca 60  
aggtgtgcac ggccacctca caaagaggcc tgaggagatt ggtgctggtg accaagggtca 120  
tatgttcggc tatgccactg acgagactcc cgagctcatg cccttgagcc atgtccttcc 180  
acgaagctcg gtgccaagct caccgaggtt cggagaacg ggacatgcc ttggctgaga 240  
cctgatggca agaccaagt cactgttgag tactacaatg acaaggggtgc catggttcca 300  
atccgcgtcc a 311

<210> 1750  
<211> 308  
<212> nucleic acid

<213> Glycine max

<400> 1750

gcangcacgc gtacgtaagc tcggaattcg gctcgangtt cttgcaacta aactcgggtgc 60  
tcgtctcacc gaggttcgca agaacggaac ctgcccattg ttgaggcctg atgggaagac 120  
ccaagtgact gttgagtatt acaatgacaa cggtgccatg gttccagttc gtgtccacac 180  
tgtgtttatc tccacccaac atgatgagac tgtgaccaac gacgaaattg cagtgcacctc 240  
aaggagcatg tgatcaagcc ggtgatcccg gagaagtacc ttgatgagaa gaccattttc 300  
catttgaa 308

<210> 1751

<211> 394

<212> nucleic acid

<213> Glycine max

<400> 1751

aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 60  
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 120  
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 180  
accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atggtttttcg 240  
gagagatcac aaccaaggcc aacgtggact atgagaagat tgtgcgtgac acatgcaagg 300  
aacattggnt tttgtctctg atgaatgttg gncttgatgc tgacaactgc aaggnccccc 360  
tcaaanattg gnncacaaaa ntccggaana ttgc 394

<210> 1752

<211> 326

<212> nucleic acid

<213> Glycine max

<400> 1752

cangcacgcg tacgtaagct cggaattcgg ctcgagggac cctaccaagg ttgacagaag 60  
tggtgcctat atcgtgaggc aggctgcaaa gagtgttgtg gcaaattggcc ttgccagaag 120  
gtgcattgtc caagtttcct atgccattgg tgtccctgag cccttgtcag tgtttgtgga 180  
catttatgga actgggaaga ttcctgacaa ggagattcctt caaattgtga aggagaattt 240

cgacttcaga cctggaatga tcaccattaa cttggacctt aagaggggtg gccataggtt 300  
cctcaagaca gctgottatg gacact 326

<210> 1753  
<211> 536  
<212> nucleic acid  
<213> Glycine max  
<400> 1753

gnnggngngt gnnttcntnn nnnntnacnn tttggcntgc cgtaccggtc cggaattccc 60  
gggtcgaccc acgcgtccgg caagccccac tcaaccacca cacctcttct cgttcacgct 120  
acccctttct gctcttcttc tacctttcaa gttttaaaag tataaagatg gcagagacat 180  
tcctatttac ctcagagtcg gtgaacgagg gacacctga caagctctgc gaccaaactc 240  
ccgatgctgt cctcgacgct tgccctcgagc aggaccaga cagcaaagtt gcctgcgaaa 300  
catgcaccaa aaccaacttg gtcattggtc tcggagaaat caccaccaag gccaacgttg 360  
actacgagaa gatagtgcgt gacacctgca ggaacatcgg cttcgtctca aatgatgtgg 420  
gactggatgc cgacaactgc aaggctctgt caacattgac agcagaccct gatattggtc 480  
aagggtggtc acgggcacct taccaaaaaa anctggaaga aattggggct tggnga 536

<210> 1754  
<211> 286  
<212> nucleic acid  
<213> Glycine max  
<400> 1754

cacgcgtacg taagctcgga attcggctcg agaccaaggt tgataggagt ggtgcttaca 60  
ttgtgagaca ggctgctaag agcattgtgg caagtggact agccagaagg tgcattgtgc 120  
aagtgtctta tgccattggt gtgcccagac ctttgtctgt ctttgttgac acctatggca 180  
ccgggaagat ccatgataag gagattctca acattgtgaa ggagaacttt gatttcaggc 240  
ccggtatgat ctccatcaac cttgntctca agaggggtgg gaataa 286

<210> 1755  
<211> 276  
<212> nucleic acid  
<213> Glycine max



<400> 1755

tagtgcctcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60  
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
atgcttgagg caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
ggtgatgggt ttccggagaga tcacaaccaa ggccaacgtg gactatgaga agattgtgcg 240  
tgacacatgc aggaacattg gttttgtctc tgatga 276

<210> 1756

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 1756

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag ggccaacgta gactatgaga 60  
agattgtccg tgacacatgc cgcgaaattg gattcatctc tgatgatgtt ggtcttgatg 120  
ctgacaaatg caaggtgttg gtcaacattg agcagcagag ccctgatatc gccagggtg 180  
tgacagggtc attcaccaag cggccagagg aggttggtgc tggtgaccag ggtcacatgt 240  
ttgggctatg ccatgatgaa acccctgagt acatgcccct cagccatgtc cttgcaacca 300

<210> 1757

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 1757

nngtcgcang cacgcgtacg taagctcgga attcggctcg agccctgata tcgcccaggg 60  
tgtgcacggg cacttcacca agcgcccaga ggagggttgg gctggtgacc agggtcacat 120  
gtttggctat gccactgatg aaaccctga gtacatgcc ctcagccatg tccttgcaac 180  
caaactcggg gctcgcctca ccgaggtag gaaaaatggg acctgtgctt ggctgaggcc 240  
agatggcaag acacaagtaa ctgttgagta ctacaatgac aatggtgcc tggttccagt 300  
tcgt 304

<210> 1758

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 1758

ngtcgcangc acgcgtacgt aagctcggaa ttcggctcga gctgcaaaga gcattgttgc 60  
aaatggactt gctaggaggg caattgngca agtttcctat gccattggtg tgctgagcc 120  
cttgtctgtg tttgttgaca cttatgggca ctgggaagat ccctgacaag gaaatcctca 180  
gcattgtgaa ggagagtttt gacttcaggc ctggcatgat ctccatcaac cttgatctca 240  
agaggggtgg aaatggcagg ttcttgaaga ctgctgcata tggacacttt ggagagatg 300  
accctgact 309

<210> 1759

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 1759

ncacgtcgan gcacgcgtac gtaagctcgg aattcggctc gaggagaaga ccatcttcca 60  
ccttaaccct tctggccgtt ttgtcattgg tggccctcat ggtgatgctg gtctcactgg 120  
aagaaagatc atcattgata cctatgggtg gtgggggtgct catggtggag gtgccttttc 180  
aggggaaggac cctaccaagg ttgacagaag tgggtgcctat atcgtgaggc aggctgcaaa 240  
gagtgtttgt gcaaattggcc ttgccagaag gtgcatgtcc aagtttcta tgccattggt 300  
gtccctgagc cctgtcagtg 320

<210> 1760

<211> 295

<212> nucleic acid

<213> Glycine max

<400> 1760

gtcgcangca cgcgtacgta agctcggaa tccgctcgag gcgtgacaca tgcaggaaca 60  
ttggttttgt ctctgatgat gttggtcttg atgctgacaa ctgcaaggctc ctctgcaaca 120  
ttgagcaaca gagtctgat attgctcaag gtgtgcacgg ccacctcaca aagaggcctg 180  
aggagattgg tgctggtgac caagtcata tgttcggcta tgccactgac gagactcccg 240  
agctcatgcc cttgagccat gtccttgcca cgaagctcgg tgccaagctc accgn 295

<210> 1761  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1761  
  
 ngtcntangc acgcgtacgt aagctcggaa ttcggctcga gggccaacat tgagcaacag 60  
 agcccggata tcgccagggt tgtgcacggc cacttcacca agcggccaga ggaggttggt 120  
 gctgggtgacc aggggtcacat gtttggggtat gccaccgatg aaacccccga gtacatgccc 180  
 ctcagccatg tccttgcaac caaacttggt gctcgctca cagagggttag gaagaatggc 240  
 acctgtgctt ggttgaggcc agatggtaag acacaagtaa ccgtcgagta ctacaat 297

<210> 1762  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1762  
  
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 ctgggggtgct catggtggag gtgccttttc aggggaaggac cctaccaagg ttgacagaag 120  
 tgggtgcctat attgtaaggc aggctgcaaa gagtgctctg gcaaattggcc ttgctagaag 180  
 gtgcattgtg caagtttctt atgccattgg tgtccctgag cccttgtcag tgtttgtgga 240  
 cacttatgga actgggaaga ttcttgacaa ggagattctg caaattgtga aggagaa 297

<210> 1763  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1763  
  
 angangacgt cgctgcacgc gtacgtaagc tcggaattcg gctcgaggga agaacgggac 60  
 atgcccttgg ctgagacctg atggcaagac ccaagtcact gttgagtact acaatgacaa 120  
 gggtgccatg gttccaatcc gcgtccacac tgtgctcatc tccacacagc atgatgagnc 180  
 tgtcacaaat gatgagattg cagctgatct taaagaacac gtgattaagc ctgtgattcc 240  
 tgagaagtac cttgatgaga agaccatttt ccatttgaac ctttctggca ggtttgtcat 300

tg

303

<210> 1764  
<211> 492  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1764

gggnaantct acgcgncag ttctggtcag agccattccc gggananaacc cagcgncn 60  
tacggctgcg agaaagacga cagaaggggg caacgcttg attttgaggg caaggcaaag 120  
ccccactcaa accaacacac ctctcctccg ttcacgtac cttttctgct cttttcttac 180  
ctttcaagtt ttaaaaagta taaagatggc agagacattc ctatttacct cagagtcggt 240  
gaacgagggg caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg 300  
cctcgagcag gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt 360  
catggtcttc ggagaaatca cgaccaaggc caacgttgac tacgagaaga tagtgctga 420  
cacctgcagg aacatcggtc tcgtctcaaa tgatgtggga ctggatgcgc acaactgcaa 480  
aggtcctcgt ca 492

<210> 1765  
<211> 295  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1765

acgtcgang cagcgctacg taagctcgga attcggctcg aggtgccttt tcagggaagg 60  
accctaccaa ggttgacaga agtggtgcct atatcgtgag gcaggctgca aagagtgttg 120  
tggcaaattg ccttgccaga aggtgcattg tccaagtttc ctatgccatt ggtgtccctg 180  
agcccttgtc agtgtttgtg gacacttatg gaactgggaa gattcctgac aaggagattc 240  
ttcaaattgt gaaggagaat ttcgacttca gacctggaat gatcaccatt aactt 295

<210> 1766  
<211> 290  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1766

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gaagctctgt gaccagatct 60  
ccgatgctgt gctcgatgca tgcttggagc aggaccctga cagcaagggt gcctgtgaaa 120  
cctgcaccaa gaccaacatg gtgatgggtt tcggagagat cacaaccaag gccaacgtgg 180  
actatgagaa gattgtgcgt gacacatgca ggaacattgg tttgtctct gatgatgta 240  
ntcttgatgc tgacaactgc aaggctcctg tcaacattga gcaacagagt 290

<210> 1767  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 1767

nagtcnncng aacgcgtngg taagctcggg anttcggctc gangtgctca tgggtggangn 60  
gccttttcag ggaaggaccc taccaagggt gacagaagtg gtgcctatat cgtgaggcag 120  
gctgcaaaga gtgttgtggc aaatggcctt gccagaagggt gcattgtcca agtttcctat 180  
gccattgggtg tccctgagcc cttgtcagtg tttgtggaca cttatggaac tgggaagatt 240  
cctgacaagg agattcttca aattgtgaag gagaatttcg acttcagacc tggaatgatc 300

<210> 1768  
<211> 327  
<212> nucleic acid  
<213> Glycine max  
<400> 1768

gcagtgtacg tnagctcgga attcggctcg agcattgggt tgcctgagcc cttgtctgtg 60  
tttgttgaca cttatggcac tgggaagatc cctgacaagg aaatcctcag cattgtgaag 120  
gagagttttg acttcaggcc tggcatgatc tccatcaacc ttgatctcaa gaggggtgga 180  
aatggcaggt tcttgaagac tgcngcatat ggacactttg gcagagatga ccctgacttc 240  
acatgggaag tggatgaagc actcaagggg gaaaaggtag tgcttaacta aaaggggttc 300  
caacactctt ggccaangga ttttgcc 327

<210> 1769  
<211> 322  
<212> nucleic acid  
<213> Glycine max

<400> 1769

gcgtacgnaa gctcgggaatt cggctcgaga atggacttgc taggagggca attgtgcagt 60  
ttcctatgcc attggtgtgc ctgagccctt gtctgtgttt gttgacactt atggcactgg 120  
gaagatccct gacaaggaaa tcctcagcat tgtgaaggag agttttgact tcaggcctgg 180  
catgatctcc atnaaccttg atctcaagag ggggtggaaat ggcaggttct tgaagactgc 240  
tgcatatgga cactttggca gagatgacct tgacttcaca tgggaagtgg tgaagccatc 300  
aagggggaga agacctgctt aa 322

<210> 1770

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 1770

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ggacatgcc ttggctgaga cctgatggca agaccaagt cactgttgag tactacaatg 120  
acaaggggtgc catggttcca atccgcgtcc aactgtgtct catctccaca cagcatgang 180  
agnctgtcac aaatgatgag attgcagctg atcttaaaga acacgnnatt aagcctgtna 240  
tncnganaa gtnccttnat gagaagacca ttttccattt gaacccttc 289

<210> 1771

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 1771

nnngcntnan gtagcgnacg taagctcgga attcggtcga aggaagaac gggacatgcc 60  
cttggctgag acctgatggc aagaccaag tcaactgttg gtactacaat gacaaggggtg 120  
ccatggttcc aatccgcgtc cacactgtgc tcatctccac acagcatgat gagactgtca 180  
caaatgatga gattgcagct gatcttaaag aacacgtgat taagcctgtg attcctgaga 240  
agtaccttga tgagaagacc attttccatt tgaacccttc tggcaggttt gtcattg 297

<210> 1772

<211> 260

<212> nucleic acid

<213> Glycine max

<400> 1772

catgcaccaa gaccaacatg gtcatgggtct ttggagagat cnntaccaag gccaacgtag 60  
actatgagaa gattgtccgt gacacatgcc gcgaaattgg attcatctct gatgatgttg 120  
gtcttgatgc tgacaatgca aggtgttggc caacattgag cagcagagcc ctgatatcgc 180  
ccaggggtgtg cacgggtcact tcaccaagcg ccagaggag gttgggtgctg gtgaccaggg 240  
tcacatgttt ggctatgcca 260

<210> 1773

<211> 338

<212> nucleic acid

<213> Glycine max

<400> 1773

ccnnccccca ccnncntacn aaagctcgga attcggtctg aggttcgggt atgccactga 60  
cgagactccc gagctcatgc ncttgagcca tgtccttgcc acgaagctnc ggtgccaaagc 120  
tcaccgaggt tcggaanaac gggacatgcc cttgggtgag acctgatggc aagaccaag 180  
tcactgttga gtactacaat gacaagggtg ccatggttcc aatccgcgtc cacactgtgc 240  
tcatctccac acagcatgat gagactgtca caaatgatga gattgcagtg atcttaaaga 300  
acacgtgatt aagcctgtga ttinctgagaa gtaccttg 338

<210> 1774

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 1774

tngnangcac gcgtacgnaa gctcggaatt cggtctgaga actttcctat tcacatctga 60  
atcagtgaac gaggggcacc ctgacaagct cctgtgacca gatctccgat gctgtgctcg 120  
atgcatgctt ggagcaggac cctgacagca aggttgcttg tgaaacctgc accaagacca 180  
acatggtgat ggttnccgga gagatcacia ccaaggccaa cgtggactat gagaagattg 240  
tgcgtgacac atgcaggaa attggttttg tctctgatga tgttggtctt gatg 294

<210> 1775

<211> 317  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1775  
  
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 gacaagctgt gcgaccagat ctctgatgca gtgctcgatg ngtgcnttga acaggaccot 120  
 gacagcaagg ttgcntgtga gacatgcacc aagaccaaca tggtcatggc ctttgagag 180  
 atcacaacca aggccaacgt agactatgag aagattgtcc gtgacacatg ccgcgaaatt 240  
 gggattcatc tctgggtggg ttggtcttga tncgtacaat gcaaggtgnt ggtcaaacat 300  
 tgagcagcag agccctg 317

<210> 1776  
 <211> 309  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1776  
  
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 cttctggcag gtttgtcagc ggagggccgc atggcgatgc tggctcacc ggccgcaaga 120  
 tcatcatcga cacctatgga ggatgggggtg cacatgggtg tggcgcttc tctgggaagg 180  
 atcctacca ggttgatagg agtgggtgcct acattgtgag gcaagctgca aagagcattg 240  
 ttgcaaattg acttgctagg aggcaattgt gcaagtttcc tatgccattg gtgtgctga 300  
 gcccttgctc 309

<210> 1777  
 <211> 329  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1777  
  
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 gttgcctgcg aaacatgcac caaaaccaac ttggtcatgg tcttncggag aaatcacgac 120  
 caaggccaac gttgactacg agaagatagt gcgtgacacc tgcaggaaca tcggcttcgt 180  
 ctcaaatgat gtgggactgg atgccgacaa ctgcaaggct ctcgtcaaca ttgagcagca 240



gagccctgat attgctcagg gtgtacacgg ccacettacc aaaaaacctg aagaaattgg 300  
 tgctggtgac cagggtcaca tgtttggct 329

<210> 1778  
 <211> 518  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1778

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 ggattccccg gtcgacccac gcgtccgtac ggctgcggaa gacgacagaa gggggcagcg 120  
 cttgatttga ggccaggcaa gcccactca accaccacac ctctcctcgt tcacgtacc 180  
 cttttctgct cttcttctac ctttcaagtt ttaaaagtat aaagatggca gagacattcc 240  
 tattttacctc agagtcggtg aacgagggac accctgacaa gctctgcgac caaatctccg 300  
 atgctgtcct cgacgcttgc ctcgagcagg acccagacag caaagttgcc tgcgaaacat 360  
 gcacaaaaac caacttggtc atggtcttcg gagaaatcac gaccaaggcc aacgttgact 420  
 acgagaagat agtgcgtgac acctgcagga acatcggtt cgtctcaaat gatgtgggac 480  
 tggatgcccg acaactgcaa ggtcctcgtt acattgac 518

<210> 1779  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1779

gtcgcangca cgcgtacgta agtcggaat tcnctcgag ntgagcaatg accaaattgc 60  
 tgctgacctt aaagagcatg ttattaagcc tgtcattcct gagaagtacc tggatgagaa 120  
 gaccatcttc caccttaacc cttctggccg ttttgtcatt ggtggccctc atggtgatgc 180  
 tggctctcact ggaagaaaga tcatcattga tacctatggt ggggtggggtg ctcatggtgg 240  
 aggtgccttt tcaggggaagg accctaccaa ggttgacaga agtgggtgcct ata 293

<210> 1780  
 <211> 269  
 <212> nucleic acid  
 <213> Glycine max

<400> 1780

cgagggacac cctgataagc tctgcgacca aatctccgat gctgtcctcg acgcttgcct 60  
cgaacaggac ccagacagca aggttgctcg cgaaacatgc accaagacca acttggtcat 120  
ggtcttcgga gagatcacca ccaaggccaa cgttgcatac gagaagatcg tgcgtgacac 180  
ctgcaggagc atcggcttca tctcanacga tgtgggactt gatgctgaca actgcaaggt 240  
ccttgtnaac attgagcagc ngagccctg 269

<210> 1781

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 1781

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagg tgggggtgctc atggtggagg 60  
tgcccttttca gggaaggacc ctaccaaggt tgacagaagt ggtgcctata tcgtgaggca 120  
ggctgcaaag agtggttggt caaatggcct tgccagaagg tgcattgtcc aagtttccta 180  
tgccattggt gtccctgagc ccttgctcagt gtttgtggac acttatggaa ctgggaagat 240  
tcctgacaag gagattcttc aaattgtgaa ggagaatttc gacttca 287

<210> 1782

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 1782

cgcngcacgc gtacgtnagc tcggaattcg gctcgaggca cggccacctc acaaagaggc 60  
ctgaggagat tgggtgctggt gaccaaggtc atatgttcgg ctatgccact gacgagactc 120  
ccgagctcat gcccttgagc catgtccttg ccacgaagct cggtgccaag ctcaccgngg 180  
ttcggaanaa cgggacatgn ccttggtgta nacctgatgg caagacncaa gtcactgttg 240  
agtactacaa tgacaagggt gccatgggtc caatccgcgt ccacactgtg ctcactctcca 300  
c 301

<210> 1783

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 1783

nagtcgcang cacgcgtacg taagctcgga attcggctcg agccagaatt catgccattg 60  
agtcattgttc ttgcaaccaa gctcgggtgct cgtctcaccg aggttcgcaa gaacggaacc 120  
tgcccatggc tgaggcctga tgggaagacc caagtgactg tggagtatta caatgataat 180  
ggtgccaggg ttccagttcg tgtncacacc gtgctaactt ccaccagca tgatgagact 240  
gtcaccaacg acgaaattgc ggctgacctc aaggagcatg tgatcaagcc tgtgatcccg 300  
gagaa 305

<210> 1784

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 1784

gtcgtgcac gcgtacgtna gctcggaatt cggctcgagt gatgatgttg gtcttgatgc 60  
tgacaaatgc aagggtgttg tcaacattga gcagcagagc cctgatatcg ccagggtgt 120  
gcacgggtcac ttcaccaagc gccagagga ggttggtgct ggtgaccagg gtcacatgtt 180  
tggctatgcc actgatgaaa ccctgagta catgcccctc agccatgtcc ttgcaaccaa 240  
actcgggtgct cgcctcaccg aggttaggaa aaatgggtacc tgtgctt 287

<210> 1785

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 1785

nancacgtcg cangcacgcg tacgtaagct cgggaattcg gctcgagagg gtcacatgtt 60  
tggctatgcc actgatgaaa ccctgagta catgcccctc agccatgtcc ttgcaaccaa 120  
actcgggtgct cgcctcaccg aggttaggaa aaatgggtacc tgtgcttggc tgaggccaga 180  
tggcaagaca caagtaactg ttgagtacta caatgacaat ggtgcatgg ttccagttcg 240  
tgtccacact gtcctaattt ccacccaaca tgatgagnct gtgagcaatn accaaattgc 300  
tgctgacctt 310

<210> 1786  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1786  
  
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 agactatgaa aagattgtcc ggcacacatg ccgcgaaatt ggattcatct ctgatgatgt 120  
 tgggtcttgat gctgacaaat gcaagggtgtt ggtcaacatt gagcaacaga gcccggatat 180  
 cggccagggt gtgcacggcc acttcaccaa gcgcccagag gaggttggtg ctggtgacca 240  
 ggggtcacatg tttgggtatg ccaccgatga aacccccgag tacatgc 287

<210> 1787  
 <211> 295  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1787  
  
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 aaggctctcg tcaacattga gcaacagagt cctgatattg ctcaagggtgt gcacggccac 120  
 ctcacaaaga ggcttgagga gattggtgct ggtgaccaag gtcatatgtt cggctatgcc 180  
 actgaagaga ctcccgagct catgcccttg agccatgtcc ttgccacgaa gctcgggtgcc 240  
 aagctcaccg aggttcggaa gaacgggaca tgcccttggc tgagacctga tggca 295

<210> 1788  
 <211> 321  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1788  
  
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 tttacctcag antcgggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat 120  
 gctgtctctg acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc 180  
 accaaaacca acttggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac 240  
 gagaagatag tgcgtgacac ctgcaggaac atcggcttcg tctcaaatga ngtgggactg 300

gatgccgaca actgcaagtc t 321

<210> 1789  
 <211> 270  
 <212> nucleic acid  
 <213> Glycine max

<400> 1789

tagtgccctcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60  
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 atgcttgagg caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
 ggtgatgggt ttccggagaga tcacaaccaa ggccaacgtg gactatgaga agattgtgcg 240  
 tgacacatgc aggaacattg gttttgtctc 270

628

<210> 1790  
 <211> 333  
 <212> nucleic acid  
 <213> Glycine max

<400> 1790

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 ggtcacatgt ttgggctatg cactgatga aaccctgag tacatgcccc tcagccatgt 120  
 cttgcaacca aactcgggtgc tcnctcacc gaggttagga aaaatggtac ctgtgcttgg 180  
 ctgaggccag atggcaagac acaagtaact gttgagtact acaatgacaa tggcgccatg 240  
 gttccagttc gtgtccacat gtcctaattt ccaccaaca tgatgagcct gtgagcaatg 300  
 accaaattgc tgctgacctt aaagagcatg tta 333

<210> 1791  
 <211> 267  
 <212> nucleic acid  
 <213> Glycine max

<400> 1791

caatggtgcc atggttccag ttcgtgtcca cactgtccta atttccaccc aacatgatga 60  
 nacctgtgag caatgaccaa attgctgctg acctaaaga gcatgttatc aagcctgtca 120  
 ttcctgagaa gtacctggat gagaagacca tcttccacct taacccttct ggccgttttg 180

tcattggtgg ccctcatggt gatgctggtc tcaactggaag aaagatcatc attgatacct 240  
atggtgggtg ggggtgctcat ggtggag 267

<210> 1792  
<211> 314  
<212> nucleic acid  
<213> Glycine max  
<400> 1792

ccanaatcgc atgcacgcgt acgtaagctc ggaattcngc tcgagctcga gccgctcgag 60  
ccggaatcag tgaacgaggg gcaccctgac aagctctgtg accagatcct ccgatgctgt 120  
gctcgatgca tgcttgagc aggaccctga cagcaagggt gctgtgaaa cctgcaccaa 180  
gaccaacatg gtgatggttt tcggagagat cacaaccaag gccaacgtgg actatgagaa 240  
gattgtgcgt gacacatgca ggaacattgg ttttgtctct gatgatgttn gtcttgatgc 300  
tgacaactgc aagt 314

<210> 1793  
<211> 512  
<212> nucleic acid  
<213> Glycine max  
<400> 1793

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nccnntgcga gaagacgaca gaagggggca acgcttnagc agacttnaca acancacaaa 120  
gcnngttact gtctgttcaa gctaacatct ccctctctct ttccttaant gcctccttnc 180  
caagaaagtt aaaatggccc aagaaacttt cctattcaca tctgaatcaa gttaacgaag 240  
gggcaccccc gacaagctct gtgaccaaga tctccgatgc tgtgctcgat gcatgcttgg 300  
agcaagacct tgacagcaan gttgcctgtg aaacctgcac caagaccaac atggtgatgg 360  
ttttcggaga gattacaacc aangccaacg tggactatga gaagattgtg cgttacacat 420  
gcangaacat tggttttgtc tctgaagatg ttggctctga agctgacaac tgcaangtcc 480  
tcgtcaacaa ttaacaacaa naattctgat at 512

<210> 1794  
<211> 294  
<212> nucleic acid

<213> Glycine max

<400> 1794

nnngntctan gcacgcgtac gtaagctcgg aattcggtc gaggaccatc ttccacctta 60  
accottcttg ccgttttctc attggtggcc ctcatggtga tgctggtctc actggaagaa 120  
agatcatcat tgatacctat ggtggctggg gtgctcatgg tggaggtgcc ttttcaggga 180  
aggaccctac caaggttgac agaagtggg cctatatgtt aaggcaggct gcaaagagtg 240  
tcgtggcaaa tggccttgct agaaggtgca ttgtgcaagt ttcctatgcc attg 294

<210> 1795

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 1795

cgannacgtc gcangcacgc gtacgtaagc tcggaattcg gctcgaggga cgagacccca 60  
gaattgatgc cattgagtca tgttcttgca actaaactcg gtgctcgtct caccgagggt 120  
cgcaagaacg gaacctgccc atggttgagg cctgatggga agaccaagt gactgttgag 180  
tattacaatg acaacggtgc catggttcca gttcgtgtcc aactgtgtct tatctccacc 240  
caacatgatg agactgtgac caacgacgaa attgcagctg acctcaagga gcatgtgatc 300  
a 301

<210> 1796

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 1796

gcatgcacgc gtacgtaagc tcggaattcg gctcgagagt ggtgcctaca ttgtgaggca 60  
agctgcaaag agcattgttg caaatggact tgctaggagg gcaattgtgc aagtttccta 120  
tgccattggt gtgcctgagc ccttgtctgt gtttgttgac acttatggca ctgggaagat 180  
ccctgacaag gaaatcctca gcattgtgaa ggagagtttt gacttcaggc ctggcatgat 240  
ctccatcaac cttgatctca agaggggtgg aaatggc 277

<210> 1797

<211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1797  
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 ggngcctata tcgtgaggca ggtgcaaaag agtggtgtgg canatggcct tgccagaagg 120  
 tgcattgtcc aagtttccta tgccattggt gtccctgagc ccttgtcagt gtttgtggac 180  
 acttatggaa ctgggaagat tcctgncaag gagattcttc aaattgtgaa ggagaatttc 240  
 gacttnagac ctggaatgat caccattaac ttggacctta agaggggtgg ccatagg 297

<210> 1798  
 <211> 264  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1798  
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 gtgctggtga ccagggtcac atgtttgggt atgncaccga tgaaaccccc gagtacatgc 120  
 ccctcagcca tgtccttgca accaaacttg gtgctgcct cagagagggt aggaagaatg 180  
 gcacctgtgc ttggttgagg ccagatggta agacacaagt aaccgtcgag tactacaatg 240  
 acaatggtgc catggttcca ttcg 264

<210> 1799  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1799  
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 tgcccatggc tgaggcctga tgggaagacc caagtgactg tggagtatta caatgataat 180  
 ggtgccaggg ttccagttcg tgtncacacc gtgctaactc ccaccagca tgatgagact 240  
 gtcaccaacg acgaaattgc ggctgacctc aaggagcatg tgatnaagcc tgtgatcccg 300  
 gngaagtnct t 311



<210> 1800  
 <211> 508  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1800  
  
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 tcacgctacc cttttctgct cttctttctac ctttcaagtt ttaaaagtat aaagatggca 180  
 gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa gctctgcgac 240  
 caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag caaagttgcc 300  
 tgcgaaacat gcacaaaaac caacttggtc atggtcttcg gagaaatcac gaccaaggcc 360  
 aacgttgact acganaagat agtgcgtgac anctgcaaga acatcggctt cntctcaaatt 420  
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 ctgatattgc ncaaggggtt naaccggc 508

<210> 1801  
 <211> 292  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1801  
  
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 gtggtgcta tatcgtgagg caggctgcaa agagtgttgt ggcaaattggc cttgccagaa 120  
 ggtgcattgt ccaagtttcc tatgccattg gtgtccctga gcccttgta gtgtttgtgg 180  
 acaattatgg aactgggaag attcctgaca aggagattct tcaaattgtg aaggagaatt 240  
 togacttcag acctggaatg atcaccatta acttggaact taagaggggt gg 292

<210> 1802  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1802  
  
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aggttgggtgc tggtgaccag ggtcacatgt ttggctatgc cactgatgaa acccctgagt 120  
 acatgcccct cagccatgtc cttgcaacca aactcgggtgc tcgcctcacc gaggttagga 180  
 aaaatgggtac ctgtgcttgg ctgaggccag atggcaagac acaagtaact gttgagtact 240  
 acaatgacaa tggtgccatg gttccagttc gtgtccacac tgtcctaatt tccaccaaac 300  
 atgatac 306

<210> 1803  
 <211> 309  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1803

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 acatggtgat ggttttcgga gagatcacia ccaaggccaa cgtggactat gagaagattg 180  
 tgcgtgacac atgcaggaac attggtttta nctctgatga tgttggtctt gatgctgaca 240  
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 gccacctca 309

<210> 1804  
 <211> 437  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1804

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 agtataaaga tggcagagan attctatatt acctcagagt cggatgaacga gggacaccct 180  
 gacaagctct ggcaccaaatt ctccgatgct gtccctcgacg cttgcctcna gcaggancca 240  
 nacagcaaaa ttgcctgcna aacatgcacc aaaaccaact tggatcatggt cttcggagan 300  
 atcacgacca aggccaacgt tgactacgag aagatagtg cgtgacacctg caggaacatc 360  
 ggcttcgtct caaaatgatg tgggactgga tcccgacaac tgcaangtcc tcgtcaacat 420  
 ttgaacanca naggcct 437

<210> 1805  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1805  
  
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 caagggtgtgc acggccacct tcacaaagag gcctgaggag attggtgctg gtgaccaagg 120  
 tcatatgttc ggctatgcca ctgacgagac tcccgagctc atgcccttga gccatgtcct 180  
 tgccacgaag ctcggtgcca agctcaccga ggttcggaag aacgggacat gcccttggct 240  
 gagacctgat ggcaagaccc aagtcactgt tgagtactac aatgacaagg gtgccatgg 299

<210> 1806  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1806  
  
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 gggactggat gccgacaact gcaaggteet cgtcaacatt gagcagcaga gccctgatat 180  
 tgctcagggt gtacacggcc accttaccaa aaaacctgaa gaaattgggtg ctggtgacca 240  
 ggggtcacatg ttgggtatg cactgatga aaccttgaa ttgatgccat tgagcc 296

<210> 1807  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1807  
  
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 gccctcatgg tgatgctggt ctactggaa gaaagatcat cattgatacc tatggtggct 180  
 ggggtgctca tggtggagggt gccttttcag ggaaggaccc taccaagggt gacagaagtg 240  
 gtgcctatat tgtaaggcag gctgcaaaga gtgtcgtggc aaatggcctt gctagaagggt 300

gcatgtgc

308

<210> 1808  
<211> 261  
<212> nucleic acid  
<213> Glycine max  
<400> 1808

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tatgccattg gtgtgcctga gcctttgtct gtgtttgttg acacctatgg cactgggaag 120  
atccatgata aggagattct caacattgtg aaggaaaact ttgatttcag gcctgggtatg 180  
atctccatca accttgatct caagaggggt ggaaataacn ggttttggn nactgccncc 240  
natggacant tggaangnac c 261

<210> 1809  
<211> 275  
<212> nucleic acid  
<213> Glycine max  
<400> 1809

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tgttgacact tatggcaactg ggaagatccc tgacaaggaa atcctcagca ttgtgaagga 120  
gagttttgac ttcaggcctg gcatgatctc catcaacctt gatctcaaga ggggtggaaa 180  
tggcagggttc ttgaagactg ctgcatatgg acactttggc agagatgacc ctgacttcac 240  
atgggaagtg gtgaagccac tcaaggggga gaagg 275

<210> 1810  
<211> 270  
<212> nucleic acid  
<213> Glycine max  
<400> 1810

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ccaacgtgga ctatgagaag attgtgcgtg acacatgcag gaacattggt tttgtctctg 120  
atgatgttgg tcttgatgct gacaactgca aggtcctcgt caacattgag caacagagtc 180  
ctgatattgc tcaaggtgtg cacggccacc tcacaaagag gcctgaggag attggtgctg 240

gtgaccaagg tcatatgttc ggctatgcc

270

<210> 1811  
<211> 317  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1811

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tgtgaggcaa gctgcaaaga gcattgttnc caaatggact tgctaggagg gcaattgtgc 180  
aagtttctta tgccattggt gtgcctgagc cttgtctgt gtttgttgac acttatggca 240  
ctgggaagat ccctgacaag gaaatcctca gcattgtgaa ggagagtttt gactcaggcc 300  
tggatgatct cnatcac 317

<210> 1812  
<211> 323  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1812

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ganacotttc tattcacatc tgaatctgta aacgagggtc accccgacaa gctgtgcgaa 120  
ccagatctct gatgcagtgc tcgatgcgtg cttgaacag gaccctgaca gcaaggttgc 180  
ctgtgagaca tgcaccaaga ccaacatggt catggtcttt ggagngatca canccaaggg 240  
ccnnacgtag nctatgagaa gattgtccgt gacacctgcc gcgaaattgg attcatctct 300  
gatgtgttcg gtcnngatgc gcc 323

<210> 1813  
<211> 342  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1813

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cctgacaagg agattcttca aattgtgaag gagaatttcg acttcagacc tggaatgatc 180  
 accattaact tggacottaa gaggggtggc cataggttcc tcaagacagc tgcttatgga 240  
 cactttggaa gggatgacct gacttcacct gggaagttgt gaagccatca agtctgagaa 300  
 gccncaactt agatgtgtga gttaaccatc ccttcattggn gc 342

<210> 1814  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1814

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 gaaggggaag aaagatcatc attgatacct atggtggctg ggggtgctcat ggtggagggtg 120  
 ccttttcagg gaaggacct accaaggttg acagaagtgg tgcctatatt gtaaggcagg 180  
 ctgcaaagag tgtcgtggca aatggccttg ctagaagggtg catttgtgtca gtttcctatg 240  
 ccattgggtgt ccctgagccc ttgtcagtgt ttgtggacac ttatgggaact gggaagattc 300  
 ctgacaagga gattctgc 318

<210> 1815  
 <211> 280  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1815

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 catgggtgggtg gtgccttctc tgggaaggat cctaccaagg ttgataggag tgggtgcctac 120  
 attgtgagggc aagctgcaaa gagcattgtt gcaaattggac ttgctaggag ggcaattgtg 180  
 caagtttctt atgccattgg tgtgcctgag cccttgtctg tgtttgttga cacttatggc 240  
 actgggaaga tccctgacaa ggaaatcctc agcattgtgn 280

<210> 1816  
 <211> 236  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1816

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 ggagattctc aacattgtga aggaaaactt tgatttcagg cctgggtatga tctccatcaa 180  
 ccttgatctc aagaggggtg gaaataacag gtttttgaag actgctgcct atggac 236

<210> 1817  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1817

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 ttgcaaattg acttgctagg agggcaattg tncaagtttc ctatgccatt ggtgngcctg 180  
 agcccttntc tgtgtttgtt gacacggatg gcaactgggaa gatccctgac aangnaatcc 240  
 tcagcattgt gaaggagagt ttgacttca ggcctggcct gatctccatc naccttgagc 300  
 tcaagagggg tgnn 314

<210> 1818  
 <211> 267  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1818

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 ttttccactt gaacccttca ggccgttttg tcattggtgg ccctcatggc gatgctggtc 180  
 tcaccggccg caagatcatt atcgatactt atggaggatg ggtgctcat ggtggtggtg 240  
 ctttctccgg gaaggaccct accaagg 267

<210> 1819  
 <211> 278  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1819

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 aggttgatag gagtgggtgcc tacattgtga ggcaagctgc aaagagcatt gttgcaaagt 120  
 gacttgctag gagggcaatt gtgcaagttt cctatgccat tgggtgtgct gagcccttgt 180  
 ctgtgtttgt tgacacttat ggcaactgga agatccctga caaggaaatc ctcagcattg 240  
 tgaaggagag ttttgacttc aggcctggca tgatctcc 278

<210> 1820  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max

<400> 1820

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 gccattggtg tgcctgagcc cttgtctgtg tttgttgaca cttatggcac tgggaagatc 120  
 cctgcacaag gaaatcctca gcattgtgaa ggagagtttt gacttcaggc ctggcatgat 180  
 ctccatcaac cttgatctca agaggggtgg aaatggcagg ttcttgaaga ctgctgcata 240  
 tggacacttt ggcagagatg accctgactt cacatgggaa g 281

<210> 1821  
 <211> 255  
 <212> nucleic acid  
 <213> Glycine max

<400> 1821

cttgccagaa ggtgcattgt gcaagtgtct tatgccattg gtgtncctga gcctttgtac 60  
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 gaaggagaac tttgatttca ggccctggtat gatctccatc aaccttgatc tcaagagggg 180  
 nggaaataac aggtttttga agactgctgc atatggacac tttggaagag aggaccctgg 240  
 acttcacatg ggaag 255

<210> 1822  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 1822



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 angcaagctg caaagagcat tgttgcaa at ggacttgcta ggagggcaat tgtgcaagtt 120  
 tcctatgcc a ttggtgtgcc tgagcccttg tctgtgtttg ttgacactta tggcactggg 180  
 aagatccctg acaaggaa at cctcagcatt gtgaaggaga gttttgactt caggctggca 240  
 tgatctccat caacttgatc tcaagagggg tgggaatggc aggttcttga gatgctgcaa 300

<210> 1823  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1823

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 gccattggnc tgccctgagcc cttgtctgtg tttgttgaca cttatggcac tgggaagatc 180  
 cctgacaagg aaatccctcag cattgtgaag gagagttttg acttcaggcc tggcatgatc 240  
 tcatcaacct tgatctcaag aggggtggaa atggcaggtt ctt 283

<210> 1824  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1824

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 agggccgcat ggcgatgntg gtctcaccgg ccgcaagatc atcatcgaca cctatggagg 120  
 atgggggtgca catggtgggtg gtgccttctc tgggaaggat cctaccaagg ntgataggag 180  
 tgggtgcctac attgtgagggc aagctgcaaa gagcattggt gcaaatggac ttgctaggag 240  
 ggcaattgtg caagtttctc atggccattg gtgtgcctga gcccttgtct gtgtttgtng 300  
 acactt 306

<210> 1825  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max

<400> 1825

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agggtcacat gtttggtat gccactgatg aaaccctga gtacatgcc ctcagccatg 120  
tccttgcaac caaactcggg gctcgctca ccgagggttag gaaaaatgg acctgtgctt 180  
ggctgaggcc agatggcaag acacaagtaa ctgttgagta ctacaatgac aatggtgcca 240  
tggttcagtg tcgtgtccac antgtntaa tttccacca ncatgatcct nctgtgagca 300  
tgaccaaatt ggt 313

<210> 1826

<211> 357

<212> nucleic acid

<213> Glycine max

<400> 1826

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ccctgacaag ctctgcgacc aaatctccga tgcgtgcctc gacgcttgcc tcgagcagga 180  
cccagacagc aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 240  
gaaatcacga ccaaggccaa cggtgactac gagaagatag tgcgtgacac ctgcaggaac 300  
atcggttcg tctcaaatga tgtgggatgg atgccgacaa ctgcaaggtc ctgctca 357

<210> 1827

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 1827

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tgagcaacag agtctgata ttgctcaagg tgtgcacggc caccttcaca aagaggcctg 120  
aggagattgg tgctgggtgac caaggctcata tgttcggcta tgccactgac gagactcccg 180  
agctcatgcc cttgagccat gtccttgcca cgaagctcgg tgccaagctc accgaggttc 240  
ggaagaacgg gacatgccct tggctgagac ctgatggcaa gaccaagtc atgttgagta 300  
tacaatgaca agggtgccat 320

<210> 1828  
 <211> 282  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1828  
  
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 gattggtgct ggtgaccaag gtcatatgtt cggtatgcc actgacgaga ctcccgagct 180  
 catgcccttg agccatgtcc ttgccacgaa gtcgggtgcc aagctcaccg aggttcggaa 240  
 gaacgggaca tgcccttggc tgagacctga tggcaagacc ca 282

<210> 1829  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1829  
  
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 cacttcacca agcgcccaga ggagggttgt gctggtgacc agggtcacat gtttggctat 180  
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 ctgctcacc gaggttagga aaaatggtac tgtgcttggc tga 283

<210> 1830  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1830  
  
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 tgagcagcag agccctgata ttgctcaggg tgtacacggc caccttacca aaaaacctga 180  
 agaaattggt gctggtgacc agggtcacat gtttggctat gccactgatg aaaccctga 240  
 attgatgcc ttgagccatg ttcttgcaac aaaactcggg gtcgtctca 290

<210> 1831  
 <211> 268  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1831  
  
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 gatcacaacc aaggccaacg tggactatga gaagattgtg cgtgacacat gcaggaacat 180  
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<210> 1832  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1832  
  
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 cttgccacga agctcgggtc caagctcacc gaggttcgga agaacgggac atgcccttgg 180  
 ctgagacctg atggcaagac ccaagtcact gttgagtact acaatgacaa gggtgccatg 240  
 gttccaatcc gcgtccacac tgtgctcatc tncacacaac atacgaccng agtgtggncg 300  
 cggattggna catgg 315

<210> 1833  
 <211> 240  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1833  
  
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 caagaacgggt acctgccctt ggctgaggcc tgatgggaag acccaagtga ccgttgagta 180  
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<210> 1834  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1834  
  
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 tttcggagag atcacaacca aggccaacgt ggactatgag aagattgtgc gtgacacatg 180  
 caggaacatt ggtccaagtc totgatgatg ttggtcttga tgcgtgacaac tgcaagggtcc 240  
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<210> 1835  
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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1835  
  
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 gataggagtg gtgcttacat tgtgagacag gctgctaaga gcatgtggca agtggacttg 180  
 ccagaagggt cattgtgcaa gtgtcttatg ccattggtgt gcttgagcct ttgtctgtgt 240  
 ttgttgacac ctatggcact gggaagatcc atgataagga gattct 286

<210> 1836  
 <211> 341  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1836  
  
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 ctgtgaccaa atctctgatg ctgtcctcga cgcttgccctc gaacaggacc cagacagcaa 180  
 gggtgcctgc gaaacatgca ccaaaaccaa cttgggtcatg gtcttcggag aaatcacgac 240  
 caaggccaat gttgactacg agaagatagt gcgtgacacc tgcaggaaca tcggctttgt 300

ctcaaacgat gtgggactgg atgccgacaa tgcaaggctc t 341

<210> 1837  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max

<400> 1837

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 ggggtgcaca tgggtgggtgt gccttctctg ggaaggatcc taccaagggt gataggagtg 180  
 gtgcctacat tgtgaggcaa gctgcaaaga gcattgttgc aaatggactt gctaggaggg 240  
 caattgtgca agtttcttat gccattgggtg tgcttgagcc cttgtctgtg tttgttgaca 300  
 cttatggcac tgg 313

<210> 1838  
 <211> 276  
 <212> nucleic acid  
 <213> Glycine max

<400> 1838

nangcacgcg tacgtaagct cggaattcgg ctcgaggaca cctgcaggaa catcggcttc 60  
 gtctcaaattg atgtgggact ggatgccgac aactgcaagg tcctcgtcaa cattgagcag 120  
 cagagccctg atattgctca ggggtgtacac ggccacctta ccaaaaaacc tgaagaaatt 180  
 ggtgctgggtg accagggtca catgtttggc tatgccactg atgaaacccc tgaattgatg 240  
 ccattgagcc atgttcttgc aacaaaactc ggtgct 276

<210> 1839  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max

<400> 1839

angtcgcang cacgcgtacg taagctcgga attcggctcg agntgcacca aaaccaactt 60  
 ggtcatggtc ttcggagaaa tcacgaccaa ggccaacgtt gactacgaga agatagtgcg 120  
 tgacacctgc aggaacatcg gcttcgtctc aaatgatgtg ggactggatg ccgacaactg 180

caaggtcctc gtcaacattg agcagcagag ccttgatatt gctcagggtg tacacggcca 240  
ccttaccaaa aaacctgaag aaattggtgc tggtgaccag ggtcac 286

<210> 1840  
<211> 315  
<212> nucleic acid  
<213> Glycine max

<400> 1840

gtcgctagca cgcgtacgta agctcggaat tcggtcgag ntttctatg ccattggtgt 60  
ccttgagccc ttgncagtgt ttgtggacac ttatggaaact gggaagattc ctgacaagga 120  
gattcttcaa attgtgaagg agaatttoga cttcagacct ggaatgatca ccattaactt 180  
ggaccttaag aggggtggcc ataggttctt caagacagct gcttatggac actttggaag 240  
ggatgacctt gacttcacct gggaagttgt gaagccactc aagtctgaga agcctcaagc 300  
ntaagattgt tgtga 315

<210> 1841  
<211> 408  
<212> nucleic acid  
<213> Glycine max

<400> 1841

gagaagacga cagaaggggg cagcgcttga tttgaggcca ggcaagcccc actcaaccac 60  
cacacctctc ctggttcacg ctaccccttt ctgctcttct tctaccttc aagttttaaa 120  
agtataaaga tggcagagac attcctatth acctcagagt cggatgaacga gggacacctt 180  
gacaagctct gcgaccaaht ctccgatgct gtctctgacg cttgcctcga gcaggaccca 240  
gacagcaaag ttgcctgcga aacatgcacc aaaaccaact tggatcatggt cttcggagaa 300  
atcacgacca aggccaacgt tgactacgag aagatagtgc gttacacctg caagaacatc 360  
cgntttctct cnaatgattt ggaactggat nccaaaaatt gcaaggte 408

<210> 1842  
<211> 255  
<212> nucleic acid  
<213> Glycine max

<400> 1842

ttgataccta tgggtggctgg ggtgctcatg gtggaggtgc cttttcaggg aaggacccta 60  
ccaaggttga cagaagtggg gcctatatattg taaggcaggc tgcaaagagt gtcgtggcaa 120  
atggccttgc tagaaggtgc attgtgcaag tttcctatgc cattgggtgc cctgagccct 180  
tgtcagtgtt tgtggacact tatggaactg ggaagattcc tgacaaggag attctgcaat 240  
tgtgaaggag attcc 255

<210> 1843  
<211> 273  
<212> nucleic acid  
<213> Glycine max  
<400> 1843

tctcaagttt ttgaagtata gagatggcag agacattcct atttacctca gagtcagtga 60  
acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc 120  
tcgaacagga cccagacagc aaggttgcct gcgaaacatg caccaaaacc aacttggtca 180  
tgggtcttcgg agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca 240  
cctgcaggaa catcggtttt gtctcaaacg atg 273

<210> 1844  
<211> 272  
<212> nucleic acid  
<213> Glycine max  
<400> 1844

nacgcangcn cgcgtacgta agctcggaat tcggctcgag agtggacttg ccagaagggtg 60  
cattgtgcaa gtgtcttatg ccattgggtg gcctgagcct ttgtctgtgt ttgttgacac 120  
ctatggcact gggaagatcc atgataagga gattctcaac attgtgaagg aaaactttga 180  
tttcaggcct ggtatgatct ccatcaacct tgatctcaag aggggtggaa ataacaggtt 240  
tttgaagact gctgcctatg gacactttgg aa 272

<210> 1845  
<211> 279  
<212> nucleic acid  
<213> Glycine max  
<400> 1845



gtcgcangca cgcgtacgta agctcggaat tcggctcgag cacgaagctc ggtgccaagc 60  
tcaccgaggt tcggaagaac gggacatgcc cttggctgag acctgatggc aagacccaag 120  
tcactgttga gtactacaat gacaaggggtg ccatggtncc caatccgctg ccacactgtg 180  
ctcatctcca cacagcatga tgagactgtc acaaatgatg agattgcagc tgatcttaaa 240  
gaacacgtga ttaagcctgt gattcctgag aagtacctt 279

<210> 1846  
<211> 269  
<212> nucleic acid  
<213> Glycine max

<400> 1846

1846

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag gaccaacatg gtgatggttt 60  
tcggagagat cacaaccaag gccaacgtgg actatgagaa gattgtgcgt gacacatgca 120  
ggaacattgg ttttgtctct gatgatgttg gtcttgntgc tgacaactgc aaggtccctc 180  
gtcaacattg agcaacagag tctgatatt gctcaagggtg tgcacggcca cctcaciaag 240  
acgcctgagg agattgggtgc tggtagacca 269

<210> 1847  
<211> 439  
<212> nucleic acid  
<213> Glycine max

<400> 1847

ccacgcgtcc gtacngctgc gagaagacga cagaaggggg cagcgcttga tttgaggcca 60  
ggcaagcccc actcaaccac cacacctctc ctcggtcacg ctaccccttt ctgctcttct 120  
tctacctttc aagtttttaa agtataaaga tggcagagac attcctattt acctcagagt 180  
cgggtgaacga gggacaccct gacgagctct gcgaccaa atccgatgct gtcctcgacg 240  
cttgccctga gcaggacca gacagcaaag ttgcctgcga aacatgcacc aaaaccaact 300  
tgggtcatggt cttcgagaaa atcacgacca angncaacgt tgactacgan aaaganantg 360  
ggttanactn gcagganntc ggcttcgtct caaatgatgt gggactggat gccgacaact 420  
gcaaggtcct cgtcaacat 439

<210> 1848

<400> 1848

[illegible]

<400>	1849
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[illegible]

<400> 1850

ctatggtggg tggggtgctc atggtggagg tgccttttca gggaaggacc ctaccaaggt 240

tgacagaagt ggtgcctata tcgtga

266

<210> 1851  
<211> 272  
<212> nucleic acid  
<213> Glycine max

<400> 1851

ggtgctcatg gtggtggtgc cttctccggg aaggatccca ccaaggttga taggagtgg 60

gcttacattg tgagacaggc tgctaagagc attgtggcaa gtggactagc cagaaggtgc 120

attgtgcaag tgtcttatgc cattgggtgtg cccgagcctt tgtctgtctt tgttgacacc 180

tatggcaccg ggaagatcca tgataaggag attctcaaca ttgtgaagga gaatttgatt 240

ncaggcccgg tatgatctcc atcaaccttg at 272

<210> 1852  
<211> 305  
<212> nucleic acid  
<213> Glycine max

<400> 1852

cnncatncgt aagtnantnc nncattnggc tcgagccaac ttggtcatgg tctncggata 60

natcntgncc aaggccancg ttgnctacga gnagatagtg cgtgacacct gcaggaacat 120

cggcttcgtc tcanatgatg tgggactgga tgccgacaac tgcaaggtcc tcgtcaacat 180

tgagcagcag agccctgata ttgctcaggg tgtacacggc caccttacca aaaaacctga 240

agaaattggg gctggtgacc agggtcacat gtttggctat gccatgatga nccctgaatt 300

gatgc 305

<210> 1853  
<211> 340  
<212> nucleic acid  
<213> Glycine max

<400> 1853

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gcacaaagcg gggttactgtc 60

tgttcaagct accatctctc tctctctttc ttagtgctc cttgccagaa gttaaaatgg 120

ccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct gacaagctct 180

gtgaccagat ctccgatgct gtgctcgatg catgcttga gcaggaccct gacagcaagg 240  
 ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag atcacaacca 300  
 aggccaacgt ggactatgag aagattgtgc gtgacacatg 340

<210> 1854  
 <211> 329  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1854

agtcgcangc acgcntacgt aagctcggaa ttcggctoga gctctctgtt ctcttctacc 60  
 tctcaagtnt ttgaagtata gagatggcag agacattcct atttacctca gactcngtga 120  
 acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc 180  
 tcgaacagga cccagacagc aaggttgctt gcgaaacatg caccaaaacc aacttggtca 240  
 tggctctcgg agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca 300  
 cctgcaggaa catcggttt gtctcaaac 329

<210> 1855  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1855

tcgcatgcac gcntacgtaa gctcggatt cggtcagagn ganattggtg ctggtgacca 60  
 gggtcacatg tttggctatg cactgatga aaccctgaa ttgatgccat tgagccatgt 120  
 tcttgcaaca aaactcgggtg ctctctcac cgaggttcgc aagaacggtg cctgcccttg 180  
 gctgaggcct gatgggaaga cccaagtgc cggttgagtat tacaatgaca atggtgccag 240  
 ggttcctatt cgtgtacaca ccgtgcnaa tctccacca acacgacgag nct 293

<210> 1856  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1856

naaangtcgc atgcacgcgt acgtaagctc ggaattcggc tcgagtgcac gttgagtatt 60

acaatgacaa tgggtgccagg gttcctattc gtgtacacac cgtgctaata tccacccaac 120  
acgacgagtn ctgtcaccaa tgacgaaatt gctgctgacc tcaaagagca tgtgatcaag 180  
cctgtgatcc cagagaagta ccttgatgag aagaccattt tccaacttga acccttcagg 240  
ccgttttgtc attggtggcc ctcatggcga tgctgggtctc accggccgca agatcattat 300  
cgatac 306

<210> 1857  
<211> 294  
<212> nucleic acid  
<213> Glycine max  
<400> 1857

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gcangcaggg gtacgtnagc tcggaattcg gctcgagggg aggttggtgc tggtgaccag 60  
ggtcacatgt ttggctatgc cactgatgaa acccctgagt acatgccct cagccatgtc 120  
cttgcaacca aactcgggtgc tcgcctcacc gaggttagga aaaatggtagc tgtgcttggc 180  
tgaggccaga tggcaagaca caagtaactg ttgagtacta caatgacaat ggtgccatgg 240  
ttccagttcg tgtccacact gtcttaattt ccacacaaca tgnnnnnacc ganc 294

<210> 1858  
<211> 394  
<212> nucleic acid  
<213> Glycine max  
<400> 1858

gagaagacga cagaaggggg cagcgcttga ttgaggcca ggcaagcccc actcaaccac 60  
cacacctctc ctggttcacg ctaccccttt ctgctcttct tctacctttc aagttttaaa 120  
agtataaaga tggcagagac attcctattt acctcagagt cggtagacga gggacaccct 180  
gacaagctct gcgaccaaatt ctccgatgct gtccctcgacg cttgcctcga gcaggaccca 240  
agacagcaaa gttgcctgcg aaacatgcac caaaaccaac ttggtcatgg tcttcggaga 300  
aatcacgacc aaggccaacg ttgactacga gaagatagtg cgtgacacct gcangaacat 360  
cggcttcgtc tcaaatgatg tgggactgga tgcc 394

<210> 1859  
<211> 307  
<212> nucleic acid

<213> Glycine max

<400> 1859

cgcatgcacg cgtacgtaag ctoggaattc gggctcgagc tctgatgatg ttggtcttga 60  
tgctgacaaa tgcaagggtg ttgtcaacnt tgagcagcag agccctgata tcgcccaggg 120  
tgtgcacggt cactttcacc aagcgcccag aggagggttg tgctgggtgnc cagggtcaca 180  
tgtttggtta tgccactgat gaaacccctg agtacatgcc cctcagccat gtccttgcaa 240  
ccaaactcgg tgctcgctc accgagttag gaaaaatggt acctgtgctt ggctgaggcc 300  
agatngc 307

<210> 1860

<211> 493

<212> nucleic acid

<213> Glycine max

<400> 1860

gnaactctta ccggccangt accgttaang agccccggt cgacannacg cgtagtccg 60  
gctgcgaaga aaacgacaga agggggcagc gcttgatttg aggccaggca agccccactc 120  
aaccaccaca cctctcctcg ttcacgctac ccctttctgc tcttcttcta cctttcaagt 180  
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 240  
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 300  
gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggc catggtcttc 360  
ggagaaatca cgaccaangg caacgttgac tacnanaann aaattncntg acacctgcag 420  
gaacatcggc ttctgtctca atgatgtggg actgggatgc cgacaactgc aangtcctcg 480  
tcaacattga gca 493

<210> 1861

<211> 489

<212> nucleic acid

<213> Glycine max

<400> 1861

ggcttttnng ccngtnnaa tcttacagn caggtaccgg tacggaattc ccggctcgac 60  
ccacgcgtac gtacggctgc gagaagacga cagaaggggg cagcgcttga tttagggcca 120

ggcaagcccc actcaaccac cacacctctc ctcgttcacg ctaccccttt ctgctcttct 180  
 tctacctttc aagtttttaa agtataaaga tggcagagac attcctattt acctcagagt 240  
 cggtgaacga gggacaccct gacaagctct ggcaccaa atctcgatgct gtccctcgacg 300  
 cttgcctcga gcaggacca gacagcaaag ttgcctgcga aacatgcacc aaaaccaact 360  
 tggatcatggt cttcggagaa atcacgacca angccaacgt tgactacgaa aaagataatt 420  
 ccttaacacc tgcaggaaca tcggttcgt ctcaa atgat gtgggactgg atgccgacaa 480  
 ctgcaaggt 489

<210> 1862  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 1862  
 gtcgcatgca cgcgtacgt agctcggaat tcggctcgag cagaacatcg ntctgtctca 60  
 aatgatgtgg gactggatgc cgacaactgc aaggtacctc gtacaacatt gagcagcaga 120  
 gccctgatat tgctcagggg gtacacggcc accttaccaa aaaacctgaa gaaattgggt 180  
 ctggtgacca gggtcacatg ttgggtatg ccaactgatga aacctctgaa ttgatgccat 240  
 tgagccatgt tcttgcaaca aaactcgggt ctcgtctcac cgagggttcgc aagaacggta 300

<210> 1863  
 <211> 330  
 <212> nucleic acid  
 <213> Glycine max

<400> 1863  
 ncatgcacgc gtacgtaagc tcggaattcg gctcgagctt ctacctctca agtttttgaa 60  
 gtatagagat ggcagagaca ttcctattta cttcagagt cagtgaacga gggacaccct 120  
 gacaagctct gtgaccaa atctctgatgct gtccctcgacg cttgcctcga acaggacca 180  
 gacagcaagg ttgcctgcga aacntgcacc aaaaccaact tggatcatggt cttcggagaa 240  
 atcacgacca aggccaatgt tgactacgag aagatagtcg gtgacacctg caggaacatc 300  
 ggctttgtct caaacgatgt gggactggat 330

<210> 1864

<211> 308  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1864  
  
 cgcatgcacg cgtacgtaag ctcggaattc ggctcganct cgagccgaat tcngctcgag 60  
 ttcggtctga gcatgggtgga gntgcctttt cagggaaagn ccctaccaag gttgacagaa 120  
 gtgggtgcta tatcgtgagg caggtcgcaa agagtgttgt ggcaaattggc cttgccagaa 180  
 ggtgcattgt ccaagtttcc tatgccattg gtgtccctga gcccttgtca gtgtttgtgg 240  
 acacttatgg aactgggaag attcctgaca aggagattct tcaaattgtg aaggagaatt 300  
 cgacttca 308

<210> 1865  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1865  
  
 gtngcangcg tacgtaagct cggaattcgg ctcgaggcct tgccagaagg tgcattgtcc 60  
 aagtttccta tgccattggt gtccctgagc ccttgtcagt gtttgtggac acttatggaa 120  
 ctgggaagat tcctgacaag gagattcttc aaattgtgaa ggagaatttc gacttcagac 180  
 ctggaatgat caccattaac ttggacctta agaggggtgg ccataggttc ctcaagacag 240  
 ctgcttatgg acactttgga agggatgacc ctgacttcac ctgggaag 288

<210> 1866  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1866  
  
 gcgtacgtna gctcggaatt cggctcgagn gcaagtttcc tatgccattg gtgtccctga 60  
 gcccttgtca gtgtttgtgg acacttatgg aactgggaag attcctgaca aggagattct 120  
 gcaaattgtg aaggagaatt tcgacttcag acctggaatg atcaccatta acttggacct 180  
 taagaggggt ggtcataggt tcctcaagac agctgcttat ggacactttg gaagggatga 240  
 tgcagacttc acctgggaag ttgtgaagcc actcaagtca g 281



<210>	1867
<211>	353
<212>	nucleic acid
<213>	Glycine max

acttaacaac	agcacaaagc	gggttactgt	ctgttcaagc	taccatctct	gctctctctn	60
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gaatcagtga	acgaggggca	cctgacaag	ctctgtgacc	agatctccga	tgctgtgctc	180
gatgcatgct	tgagcaggac	cctgacagca	aggttgccctg	tgaaacctgc	accaagacna	240
acatggtgat	ggttttcgga	gagatcacia	ccaangccaa	cgtggactat	gaggagattg	300
tgngtgacac	atgcaagaac	attggtttgt	ctccgatgat	gtnggtcttn	ntg	353

<400> 1868

aatatttcacg	ccgnccaggt	ancggtcana	gaatttcccg	ggncgacca	cgcgtoconcg	60
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gcagcgcttg	agaccaagcc	ccactcaacc	accacaccac	tctctctgct	cttcttctac	180
ctttcaagtt	tttaaagtat	taagatggca	gagacattcc	tatttacctc	agagtcagtg	240
aacgagggac	accctgacaa	gctctgcgac	caaatctccg	atgctgtcct	cgacgcttgc	300
cttgaacagg	accagacag	caaggttgcc	tgcgaaacat	gcaccaaaga	ccaacttggt	360
catggtcttc	ggagagatta	acancaaagg	ccaacgttga	ctacgaagaa	gatcgtgcgt	420
gacacctgca	ggaacatcgg	ctccgtctca	aacgatgtgg	gacttgatgc	tgacaactgc	480
aangtccttg	taacaatgaa	ca				502

naangnagan gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gtttttgaag 60  
tatagagatg gcagagacat tcctatttcc ctcagagtca gtgaacgagg gacaccctga 120  
caagctctgt gaccaaactct ctgatgctgt cctcgacgct tgcctcgaac aggaccaga 180  
cagcaagggtt gcctgcgaaa catgcaccaa aaccaacttg gtcatgggtct tcggagaaat 240  
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tttgtctcna acgatgtggg at 322

<210> 1870  
<211> 418  
<212> nucleic acid  
<213> Glycine max

<400> 1870

tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 60  
aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 120  
aagctctgtg accagatctn cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 180  
agcaagggtt cctgtgaaac ctgcaccaag accaactggt tgatgggttt cggagagatc 240  
acaaccaagg ccaacgtgga ctatgagaaa gattgtgcgt gacacatgca ggaaccattg 300  
ggttttgctc tgatgaatgt ggtcttggat gcttgacact gcaaggctcct cgtcaacatt 360  
tgagcaacag aagtctgat antgnttcaa ggtgtgcacg ggcacottac aaaagang 418

<210> 1871  
<211> 261  
<212> nucleic acid  
<213> Glycine max

<400> 1871

tacctctcaa gtttttgaag tatagagatg gcagagacat tcctatttac ctcagagtca 60  
gtgaacgagg gacaccctga caagctctgt gaccaaactct ctgatgctgt cctcgacgct 120  
tgcctcgaac aggaccaga cagcaagggtt gcctgcgaaa catgcaccaa aaccaacttg 180  
gtcatgggtct tcggagaaat cacgaccaag gccaatgttg actacgagaa gatagtgcgt 240  
gacacctgca ggnacatcgg g 261

<210> 1872

<211> 277  
 <212> nucleic acid  
 <213> Glycine max

<400> 1872

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acgtcgcacg cacgcgtacg taagctcgga attcggctcg agcgaggggc accctgnaca 60
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 120
gcaagggttc ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca 180
caaccaaggc caacgtggac tatgagaaga ttgtgcgtga cacatgcagg aacattgggt 240
ttgtctctga tgatgttggt cttgatgctg acnactg 277
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<210> 1873  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max

<400> 1873

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ctgnacgnnt gctcggaatt cggctcgaga ccaaggccaa cgttgactac gagaagatag 60
tgctgacac ctgcaggaac atcggttcg tctcaaatga tgtgggactg gatgccgaca 120
actgcaaggt cctcgtcaac attgagcann cagagccctg atattgctca ggggtgtacac 180
ggccacctta ccaaaaaacc tgaagaaatt ggtgctggtg accaggggtca catgtttggc 240
tatgccactg atgaaacccc tgattgatgc cattgagcca tgttcttgca a 291
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<210> 1874  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max

<400> 1874

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agtgaacgag gggcacctcg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120
atgcttgagg caggaccctg acagcaaggt tgctgtgaaa cctgcaccaa gaccaacatg 180
gtgatggttt tcggagagat cacaaccaag gccaacgtgg atatgagaag atgtgcgtga 240
cacatgcagg aacattgggt ttgctctgat gaggttggctt gatgctg 287
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<210> 1875

<211> 262  
 <212> nucleic acid  
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 <400> 1875  
  
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 cctacattgt gaggcaagct gcaaagagca ttgttgcaaa tggacttgct aggagggcaa 120  
 ttgtgcaagt ttctatgcc attggtgtgc ctgagccctg tctgtgtttg ttgacactta 180  
 tggcactggg aagatccctg acaaggaaat cctcagcatt gtgaaggaga gttttgactt 240  
 caggcctggc atgatctcca tc 262

<210> 1876  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1876  
  
 tcncangcac gcgtacgtaa gctcggaatt cggctcgagg gggnanctga caagctctgt 60  
 gancagatct ccgatgctgt gctcgatggc atgcttgag caggacctt acagcaangt 120  
 tgcctgtgan acctgnanca agaccaacat ggtgatgggt tncngagaga tcanaaccaa 180  
 ggccaacgtg gactatgaga agattgtgcg tgacacatgc aggaacattg gttttgtctc 240  
 tgatgatggt ggtctgatgc tgacaatgca nagtcctcgt caacattgag cnacagagtc 300  
 ctgatattgc tcaag 315

<210> 1877  
 <211> 489  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1877  
  
 ggnngnntnn nggnaaactt tcaacttgccg cgnccaggta anggtcagga attcccgggt 60  
 cgaccacgc gtccgtacgg ctgcgagaag acgacagaag ggggcagcgc ttnagatnan 120  
 nccccactca accaccacac cactctctct gctcttcttc tacctttcaa gtttttaaag 180  
 tattaagatg gcagagacat tcctatttac ctgagagtca gtgaacgagg gacaccctga 240  
 caagctctgc gaccanactt ccgatgctgt cctcgacgct tgccttgaac angaccaga 300

cagcaagggtt gcctgcgaaa catgcaccaa naccaacttg gtcatggtct tcggagaaga 360  
tnancaccaa ggccaacggt gactacgaag aagatcgtgc gttgacacct gcangaacat 420  
cggcttccgt ctcaaacgat gtgggacttg atgctgacaa ctgcaanggt cttgtnaaca 480  
ttgancacc 489

<210> 1878  
<211> 468  
<212> nucleic acid  
<213> Glycine max  
<400> 1878

actttacgnt gccangtccc ggtcangaat ncccggtcg acccacgcgt cngacggctg 60  
cgagaagacg acagaagggg gcagcgcttg agaccaagcc ccaactcaacc accacaccac 120  
tctctctgct cttctttctac ctttcaagtt tttaaagtat taagatggca gagacattcc 180  
tatttacctc agagtcagtg aacgagggac accctgacaa gctctgacgac caaatctccg 240  
atgctgtcct cgacgcttgc cttgaacagg acccagacag caaggttgcc tgcgaaacat 300  
gcaccaagac caacttggtc atggtcttcg gagaagatna ccaccaaggc caacgttgac 360  
tacgaagaag atngtgcgtg acacctgcan gaacatcngc ttcgtctcaa aagaatttgg 420  
acttgatcct gaacaactgc aaaggtccct tgtaaacaat naancacc 468

<210> 1879  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 1879

acgtcgcang cacgcgtacg taagctcgga attcggctcg agccccgag tacatgcccc 60  
tcagccatgt ccttgcaacc aaacttggtg ctgcctcac agaggtagg aagaatggca 120  
cctgtgcttg gttgaggcca gatggaaga cacaagtaac cgtcgagtac tacaatgaca 180  
atggtgccat ggttccagtt cgtgtccaca ctgtcctaatt ttccacccaa catgatgnga 240  
cgtgagcaat gatcaaattg ctgcggacct taaagagcat gttatcaagc ctgtcattcc 300

<210> 1880  
<211> 477  
<212> nucleic acid

<213> Glycine max  
 <400> 1880  
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 tacggctgcg agaagacgac agaagggggc agcgcttgat ttgaggccag gcaagcccca 120  
 ctcaaccacc acacctctcc tcgttcacgc tacccttttc tgctcttctt ctacctttca 180  
 agttttaaaa gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag 240  
 ggacaccctg acaagctctg cgaccaaata tccgatgctg tctctgacgc ttgcctcgag 300  
 caggaccag acagcaaagt tgcttgcgaa acatgcacca aaaccaactt ggtcatggtc 360  
 ttcggagaaa tcacgaccaa ggccaacggt gactacgaga agatagtgcg tgacacctgc 420  
 aagaacaacc ggttttctccc naattgaatn tgggactgga tgccgacaac tgcaang 477

<210> 1881  
 <211> 259  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1881

tgcccttggc tgagacctga tggcaagacc caagtcactg ttgagtacta caatgacaag 60  
 ggtgccatgg ttccaatccg cgtccacact gtgctcatct ccacacagca tgatgagnct 120  
 gtcacaaatg ntgagattgc agctgatctt aaagaacacg tgattaagcc tgtgattcct 180  
 gagaagtacc ttgatgagag accattttcca tttgaaccct tccnggcagg ttgtcattgg 240  
 agggcgggcat ggggatgng 259

<210> 1882  
 <211> 254  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1882

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 tgttcttgcn actnanctcg gtgctcgtct caccgagggt cgnagaacg gaacctgccc 120  
 ntggttgagg cctgatggga agaccaagt gactgttgag tattacnntg acaacgggtgc 180  
 catngttcca gttcgtgtcc nactgtgct tatctccacc caacntgatg ngntgngacc 240

aacgacgaaa ttgc

254

<210> 1883  
<211> 279  
<212> nucleic acid  
<213> Glycine max

<400> 1883

nanactnaat gcacgcgtac gtaagctcgg aattcggctc gagagacagc aaagttgcct 60  
gcgaaacatg caccaaaaacc aacttgggtca tggctcttcgg agaaatcacg accaaggcca 120  
acgttgacta cgagaagata gtgcgtgaca cctgcaggaa catcggcttc gtctcaaatg 180  
atgtgggact ggatgccgac aactgcaagg tctcgtcaa cattgagcag cagagccctg 240  
atattgctca ggggtgtacac ggccacctta ccaaaaaaac 279

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<400> 1884

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ctcttctacc tctcaagttt ttgaagtata gagatggcag agacattcct atttacctca 120  
gagtcagtga acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc 180  
gacgcttgcc tcgaacagga cccagacagc aaggttgcct gcgaaacatg caccaaaaacc 240  
aacttgggtca tggctcttcgg agaaatcacg accaaggcca atgttgacta cgagaagata 300  
gtgcgtgaca cct 313

<210> 1885  
<211> 299  
<212> nucleic acid  
<213> Glycine max

<400> 1885

acgtcgcgatg cacgcgtacg tnagctcgga attcggctcg agcttctacc tctcaagttt 60  
ttgaagtata gagatggcag agacattcct atttacctca gagtcagtga acgagggaca 120  
ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc tcgaacagga 180

cccagacagc aaggttgctt gcgaaacatg caccaaaacc aacttggtca tggctcttcgg 240  
 agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca cctgcagga 299

<210> 1886  
 <211> 301  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1886

nntaannntn agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gaaagtataa 60  
 agatggcaga gacattccta ttacctcag agtcggtgaa cgaggacac cctgacaagc 120  
 tctgcgacca natctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca 180  
 aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaaatcacga 240  
 ccaaggccaa cgttgactac gagaagatag tgcgtgacac ctgcaggaac atcggtctcg 300  
 t 301

<210> 1887  
 <211> 508  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1887

tnnnnnaact nagttctccg cngcctgna angtaagan gncccgggcc gacccacgcg 60  
 tccntacggc tgcgagaaga cgacagaagg gggcagcgt tgatttgagg ccaggcaagc 120  
 cccactcaac caccacacct ctctcgttc acgtacccc tttctgctct tttctacct 180  
 ttcaagtttt aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa 240  
 cgaggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt 300  
 cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat 360  
 ggtcttcgga gaaatcacga ccaaggccaa cgttgactac gagaagatag tgcgtgacac 420  
 ctgcangaac atcggtctcg tctcaaaatg atgtnggact ggattccac aactgcaaag 480  
 ncctccnta aaatttggc anaancc 508

<210> 1888  
 <211> 278  
 <212> nucleic acid



<213> Glycine max

<400> 1888

gncgcacgcg tacgtaagct cggaattcgg ctcgagggca ggtttgtcat tggagggccg 60  
catggcgatg ctggtctcac cgcccgcaag atcatcatcg acacctatgg aggatggggg 120  
gcacatgggtg gtgggtgncct tctctgggaa ggatcctacc aagggttgnta ggagtgggtgc 180  
ctacattntg aggcaagctg caaagagcat tgttgcaaat ggacttgcta ggagggcaat 240  
tgtgcaagtt tcctatgccca ttgggtgtgcc tgagccct 278

<210> 1889

<211> 280

<212> nucleic acid

<213> Glycine max

<400> 1889

cnttggtga gacctgatgg caagacccaa gtcantgttg agtactacaa tgacaagggt 60  
gccatggttc caatnccggt ccacantgtg cttcatntnc acacagcatg atgagtgngt 120  
nanaaatgat gagattgcag ctgatcttaa agaacacgtg attaagcntg tgattnctga 180  
gaagtacctt gatgagaaga ccattttcca tttgaacct tntgggcagg tttgtcatgg 240  
agggccgcag ggcgattttg gtgtnanggc ngnaagatcc 280

<210> 1890

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 1890

ncncangcnc gcnnncgtga gctcgnatt cggctcgagg tttttgangt atagagatgg 60  
cagagacatt cctatttacc tcagagtcag tgaacgaggg acacctgan aagctctgtg 120  
accaaattctc tgatgctgtc ctgacgctt gcctcgnaca ggaccagac agcaagggtg 180  
cntgcgaaac atgcacaaaa accaacttgg tgcatgggtct tcggagaaat cacgaccaag 240  
gccaatgttg actacgagaa gatagtgcgt gacacctgca ggaacatcgg ctttgtctca 300  
aacgatgtgg 310

<210> 1891

<211> 290  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1891  
  
 cagcangcac gcgtacgtaa gctcggaatt cggctcgaga ccaacttggn atggtcttcg 60  
 gagaaatcac gaccaaggcc aacgttgact acgagaagat agtgcgtagac acctgcagga 120  
 acantcggct tcgtctcaaa tgatgtggga ctggatgccg acaactgcaa ggtcctcgtc 180  
 aacattgagc agccgagccc tgatattgct caggggtgtac acggccacct taccaaaaaa 240  
 cctgaagaaa ttgggtgctgg tgaccagggt cacatgtttg gctatgccat 290

<210> 1892  
 <211> 502  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1892  
  
 tttaaactct ccgcggtcag gtaacggctn gngaattccc gggncgaccc angcgtccan 60  
 nccaggcgtc agcccaacgc gtongtgngg ctgcnagaag annacanaag ggggcagcgc 120  
 ttgatttnag gccaggcang cccactcat ccancanacc tctcctcgtt cangtncct 180  
 ctttctgccc ttontctacc tttcangttt taaaagtata nagatggcag agacattcct 240  
 atttacctca gagtcggtga acgagggaca ccctgacaag ctctgctacc aaatctccga 300  
 tgctgtcctc gacgcttgcc tcgagcanga ccagacagc naagttgcct gcgaaacatn 360  
 caccatancc aacttggtca tgggtcttcgg aganatcacg accaaggcca acgttgacta 420  
 cgaagaagat agtgcgtagac acctgcagga acatcngntt cgtctcaaata tatgtgggac 480  
 tggatgccan canctgcaag gt 502

<210> 1893  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1893  
  
 tcgcntgcac gcgtacgtaa gctcggaatt cggctcgagn aagttgcctg cganacatgc 60  
 accaanacca acttggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac 120

gagaagatag tgcgtgacac ctgcaggaac atcggttcg tctcaaata tgtgggactg 180  
 gatgccgaca actgcaaggt cctcgtcaac attgagcngc agagccctga tattgctcag 240  
 ggtgtacacg gccaccttac caaaaaacct gaagaaannc ntgctg 286

<210> 1894  
 <211> 326  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1894

ncgcngcacg cgtacgtaag ctcggaattc ggctcgagca caaagcgggt tactgtctgt 60  
 tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt aaaatggccc 120  
 aagaaaacttt cctattcaca totgaatcag tgaacgaggg gcaccctgac aagctctgtg 180  
 accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac agcaagggtg 240  
 cctgtgaaac ctgcaccaag accaaccatgg tgatggtttt cggagagatc acaaccaagg 300  
 ccaacgtgga ctatgagaag attgtg 326

<210> 1895  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1895

tcgnangcaa gcncngaant cngcncnagc gnnnntagnc nanngtgntt accggcngca 60  
 agatcantat cgatacttat nggaggatgn ngtgctcatg gtngtggtgc tttctccggc 120  
 aaggacccta ccaagggtga taggagtggg gcttacattg tgagacaggc tgctaaganc 180  
 attgtggcaa gtggactngc cagaagggtc attgtgcaag tgtcttatgc cattgggtgtg 240  
 cctgagcctt tgtctgtggt tgttgacacc tatggcactg ganagatccc tgacaaggag 300  
 atct 304

<210> 1896  
 <211> 273  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1896

ncgtcgcatg cacgcgtacg tnagctcgga attcggtcgc agctgtgctt ggctgaggcc 60  
 agatggcaag acacaagtaa ctgttgagta ctacaatgac aatgggtgcca tggttccagt 120  
 tcgtgtccac actgtcctaa tttccacca acatgatgac nangtgagca atgaccaa 180  
 tgctgtgac cttaaagagc atgttatcaa gctgtcatt cctgagaagt acctggatga 240  
 gaagaccatc ttccacctta acccttctgg ccg 273

<210> 1897  
 <211> 334  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1897

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag cagcacaaag cgggttactg 60  
 ncctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120  
 tggcccaaga aactttccta ttcacatctg aatcagtga cgaggggcac cctgacaagc 180  
 tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240  
 aggttgcttg tgaaacctgc accaagacca acatggtgat ggttttcgga gagatcacia 300  
 ccaaggccaa cgtggactat gagaagattg tgcg 334

<210> 1898  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1898

gtcgngcac gcgtacgtaa gctcggaatt cggctcgagn cctcacagag gttaggaaga 60  
 atggcacctg tgcttggttg aggccagatg gtaagacaca agtaaccgtc gagtactaca 120  
 atgacaatgg tgccatgggt ccagttcgtg tccacactgt cctaatttcc acccaacatg 180  
 acgacctgtg agccatgatc aaattgctgc ggaccttaaa gancatgtta tcaagcctgt 240  
 cattcctgag aagtaccttg atgagaagac catcttccac ttaacccttc tgg 293

<210> 1899  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max

<400> 1899

cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 60  
tataaagatg gcagagacat tectattttac ctacagagtcg gtgaacgagg gacaccctga 120  
caagctctgc gaccaaattct ccgatgctgt cctcgacgct tgcctcgagc aggaccacaga 180  
cancaaagtt gcctgcgaaa catgcaccaa aaccaacttg gttcatggtc ttcgagagaaa 240  
tcacgaccaa ggccaacggt gactacgaga agatagtgcg tgacacctgc aggaacatcg 300  
gcttcgtctc aaatga 316

<210> 1900

<211> 279

<212> nucleic acid

<213> Glycine max

<400> 1900

ttcctattta cctcagagtc ggtgaacgag ggacaccctg acaagctctg cgaccaaata 60  
tcgatgctg tctacgacg cttgcctcga gcaggacca gacagcaaag ttgcctgcga 120  
aacatgcacc aaaaccaact tggatcatgg cttcgagaaa atcacgacca aggccaacgt 180  
tgactacgag aagatagtc gtnacactgg agganacatg ggcttcgtct naaatgntgn 240  
gggactggat cccganaant gaaggtcncg aaaatntga 279

<210> 1901

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 1901

cgctgcacgc gtacgtaagc tcggaattcg gctcgagatg ctgtcctcga cgcttgctc 60  
gagcaggacc cagacagcaa agttgcctgc gaaacatgca ccaaaccacaa cttggatcatg 120  
gtcttcggag aaatcacgac caaggccaac gttgactacg agaagatagt gcgtgacacc 180  
tgcaggaaca tcggcttcgt ctcaaatgat gtgggactgg atgccgacaa ctgcaagggt 240  
cctcgttcaa cattgagcag cagagccctg atattgctca ggggtg 285

<210> 1902

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 1902

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag caggacccag acagcaaagt 60  
tgcttgcgaa acatgcaaaa accaacttgg tcatgggtctt cggagaaatc acgaccaagg 120  
ccaacgttga ctacgagaag atagtgcgtg acacctgcag gaacatcggc ttctgtctcaa 180  
atgatgtggg actggatgcc gacaactgca aggtcctcgt caacattgag cagcagagcc 240  
ctgatattgc tcagggtgta caccggccacc ttaccaaaaa ac 282

<210> 1903

<211> 476

<212> nucleic acid

<213> Glycine max

<400> 1903

tttactnnnc cngngccatg taanagtana gaagtcccg gccgaccac gngtcnntac 60  
ggctgcgaga agacgacaga agggggcagc gcttgatttg aggccaggca agccccactc 120  
aaccaccaca cctctctctg ttcacgttac cctttctgc tctttctcna cctttcaagt 180  
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 240  
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcat 300  
gaccagaca gcaaagttgc ctgcgaaaca tgcaccagaa ccaacttggc catggtcttc 360  
ggagaaatca cgaccatggg caacgttgac taccagaaga taagtgcgtg acacctgcag 420  
gaacatcggg ttctgtctcaa atgatgtggg actggatgcc nacaattctg anangg 476

<210> 1904

<211> 496

<212> nucleic acid

<213> Glycine max

<400> 1904

aactttcgtg ccagnccggt caagaatncc gggtcgaccc acgcgtccgt acggctgcga 60  
gaagacgaca gaagggtacg gctgcgagaa gacgacagag ggggcagcgc ttgatttgag 120  
gccaggcaag cccactcaa ccaccacacc tctcctcgtt cagctaccc ctttctgctc 180  
ttcttctacc tttcaagttt taaaagtata aagatggcag agacattcct atttacctca 240

gagtcggtga acgagggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc 300  
gacgcttgcc tcgagcagga cccagacagc aaagttgctt gcgaaacatg caccaaaaacc 360  
aacttggtca tggctcttcgg agaaatcacg accaaggcca acgttgacta cgaagaagat 420  
agtgcgtgac acctgcagga acatcggtt cgtctcaaat gatgtgggac tgggatgccg 480  
acaactgcaa ngtcct 496

<210> 1905  
<211> 247  
<212> nucleic acid  
<213> Glycine max  
<400> 1905

nannatcgcn tgcacgcgta cgtnagctcg gaattcggct cgagnntgga ttcattctctg 60  
atgatgttgg tcttgatgct gacaaatgca aggtgttggc caacattgag cagcagagcc 120  
ctgatatcgc ccagggtgtg cagggtcact tcaccaagcg cccagaggag gttggtgctg 180  
gtgaccaggg tcacatgttt ggctatgcc a ctgatgaaac ccctgagtac atgcccctca 240  
gccatnt 247

<210> 1906  
<211> 308  
<212> nucleic acid  
<213> Glycine max  
<400> 1906

atncacatgt cacangcacg cgtacgtaag ctcggaattc ggctcgagnc tntcncttgt 60  
ntgtgnatgc tgacacttat ggactggga agatccctga caaggaaatc ctcagnattg 120  
tgaaggagag ttttgacttc aggcctggnn tgatctccat caaccttgnt ctcaagaggg 180  
gtggaaatgg caggttcttg aagactgctg catatggnc ctttggcaga natgaccng 240  
acttcacatg ggaantggtn angcgactca aggggganna ggtaccagct tancatanaag 300  
ggntcct 308

<210> 1907  
<211> 292  
<212> nucleic acid  
<213> Glycine max

<400> 1907

ancgnactgc acgcgtacgt aagctcggaa ttcggtctga gaaagtataa agatggcaga 60  
gncattccta tttaacctag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 120  
aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcttg 180  
cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaaatcacga ccaaggccaa 240  
cgttgactac gagaagatat ngngtgacac ctgcaggaac atcggtcttg tc 292

<210> 1908

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 1908

agtcgcangc acgcgtacgt aanctcggaa ttcggtctga ggtttggcta tgccactgta 60  
tgaaaccctt gactacatgc cctcagcca tgccttgca accaaactcg gtgctgcctt 120  
caccgaggtt aggaaaaatg gtacctgtgc ttggctgagg ccagatggca agacacnagt 180  
aactgttgag tactacaatg acaatggtgc catgggtcca gttcgtgtcc aactgtcctt 240  
aatntncacc caacatgatg annngtgag caatgaccaa attgctgctg gaccttaaag 300

<210> 1909

<211> 458

<212> nucleic acid

<213> Glycine max

<400> 1909

atttcaggtg gtcttatagg ccaanaatga cgtaagacgc acgcgtncgt aanctcggaa 60  
ttcggtctga gggccanggc aagccccact caaccaccac acctctcttc gttcacgcta 120  
cccccttctg ctcttcttct acctttcaag ttttaaaagt ataaagatgg cagagacatt 180  
cctatttacc tcagagtcgg tgaacgaggg acaccctgac aagctctgcg accaaatctc 240  
cgatgctgtc ctgcagcgtt gcctcgagca ggaccanac agcaaagttg cctgcgaaaac 300  
atgcacaaaa accaacttgg ncatggtctt cgagagaaac acgaccaagg ccaacgttga 360  
ctacgagaag atagtgcgtg acacctgcag gaacatcggg ctctgtctca aatgatgtgg 420  
gactggatgc cgacaantgc aangttctcg tcaacant 458



<210>	1910
<211>	308
<212>	nucleic acid
<213>	Glycine max

<400> 1910

ngncncangc	ncgcntacgt	nanctcggaa	ttcggtctga	gctaccatct	tctctctctc	60
tttcttagtg	cctccttgcc	agangtnaaa	atggcccaag	aaacttnoct	atncacatct	120
gantcagtga	acgaggggca	ccttgacaag	ctctgtgacc	agatctccga	tgtctgtgctc	180
gntgcatgct	tggagcagga	ccctgacagc	aaggttgcct	gtgaaacctg	caccaagacc	240
aacatggtga	tggttttcgg	agagatcaca	accaaggcca	acgtggacta	tgagaagatt	300
gtgcgtga						308

<210>	1911
<211>	306
<212>	nucleic acid
<213>	Glycine max

<400> 1911

cgtgtacgta	agctcggaat	tcggctcgag	aagtttttga	agtatagaga	tggcagagac	60
attcctat	acctcagagt	cagtgaacga	gggacaccct	gacaagctct	gtgaccaa	120
ctctgatgct	gtcctcgacg	cttgccctoga	acaggaccca	gacagcaagg	ttgcctgcga	180
aacatgcacc	aaaaccaact	tggtcatggt	cttcggagaa	atcacgacca	aggccaatgt	240
ngactacgag	aagatagtgc	gtgacactgc	aggacatngg	tttgtccnaa	cgngnggncn	300
gncccc						306

<210>	1912
<211>	504
<212>	nucleic acid
<213>	Glycine max

<400> 1912

aatgncaaac tncncagccg nccctgncnng gtcataagggn ccgacngacc cgteocnaacc 60  
acgnatccgc tgnacantgn gcngacgcgt gggctgcnag aagacgacag aagggggcag 120  
cgcttgattt gaggccaggc aagccccact caaccaccac acctctcttc gttaacqcta 180

cccctttctg ctctttcttct acctttcaag ttttaaaagt ataaagatgg cagagacatt 240  
 cctatttacc tcagagtcgg tgaacgaggg acaccctgac aagctctgcg accaaatctc 300  
 cgatgctgtc ctcgacgctt gcctcgagca ggaccagac agcaaagttg cctgcgaaac 360  
 atgcacaaaa accaacttgg tcatggtctt cggagaaatc acgaccaang ccaacgttga 420  
 ctacgagaag atagtgentg acacctgcac ggaaatnggg ttcttctcaa ttaatttggg 480  
 acgggtttcc cnaaaactnc aagg 504

<210> 1913  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max

<400> 1913

tcnatgcacg cgtacgtaag ctcggaattc ggctcgagtc aagggtgtgca cggccacctt 60  
 cacaagagg cctgaggaga ttggtgctgg tgaccaaggt catatgttcg gctatgccac 120  
 tgacgagact cccgagctca tgcccttgag ccatgtcctt gccacgaagc cggtgccaag 180  
 ctcaccgagg ttcggaagaa cgggacatgc cttgggctga gacctgatgg caagacccaa 240  
 gtcactgttg agtactacaa tgacaagggt gccatggttc caatccgcg 289

<210> 1914  
 <211> 345  
 <212> nucleic acid  
 <213> Glycine max

<400> 1914

gnattgnagt acgcgtacgt nagctcgga ttcggctcgn ggacttaaca acaacncana 60  
 gngggttann gtctgttcaa gctaccatct ctctctctct ttcttagtgn ctctttgcna 120  
 gaagttaana tggccaaga nactttccta ttcacatctg aatcagtga cgaggggcac 180  
 cctgacaagc tctgtgacca natctccgat gctgtgctcg atgcatnctt ggagcaggac 240  
 cctgacagca aggttgctg tgaaacctgc accaanacna acatgggtgat tgttttcgga 300  
 gagatcacia ccaaggccaa cgttgactat gagaagattg tgcnt 345

<210> 1915  
 <211> 331  
 <212> nucleic acid

<213> Glycine max

<400> 1915

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60  
ggttactgtc tgttcaagct accatctctc tctctctttc ntagtgctc cttgccagaa 120  
gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
gacagcaagg ttgctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300  
atcacaacca aggccaacgt ggactatgag a 331

<210> 1916

<211> 244

<212> nucleic acid

<213> Glycine max

<400> 1916

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc caaggccaac gtagactatg 60  
aaaagattgt ccgcgacaca tgccgcgaaa ttggattcat ctctgatgat gttggctctg 120  
atgctgacaa atgcaagggtg ttggtcaaca ttgagcaaca gagcccggat atcgcccagg 180  
gtgtgcacgg ccacttcacc aagcgcccag aggaggttgg tgctgggtgac cagggtcaacn 240  
tggt 244

<210> 1917

<211> 290

<212> nucleic acid

<213> Glycine max

<400> 1917

nnngtngcat gcncgcgtac gtaagctcng nntncggctc tntgcctata tcgtgagnca 60  
ggctgcnaag agtggtgtgg naaatggcct tgccagaagg tgcattgtcc nagtttctta 120  
tnccattggt gtccttgagc ccttgctcagt gtttatggac acttatggaa ctgggaanat 180  
tcctgacaag gngattcttc aaattgtgna ggagaatttc gacttcagac ctggaatgat 240  
caccattaac ttggacctta agaggggtgg ccatagggtc ctcaagacag 290

<210> 1918

<211> 314  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1918  
  
 nnaagttnan gcacgcntac gtaagctcgg aattcggctc gaggttactg tctgttcaag 60  
 ctaccatctc tctctctott tottagtgcc tcttgccag aagttaaaat ggccaagaa 120  
 actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagct ctgtgaccag 180  
 atctccgatg ctgtgctcga tgcattgctt gagcaggacc ctgacagcaa ggttgctgt 240  
 gaaacctgca ccaagaccaa catggtgatg gttttcggag agatcacaac caaggccaac 300  
 gtggactatg agaa 314

<210> 1919  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1919  
  
 tattcnnnac gtgcgatgca cgcgtacgta agctcggaat tcggctcgag cactctctct 60  
 gctotttcttc tacctttcaa gtttttaaag tattaagatg gcagagacat tcctatttac 120  
 ctcagagtca gtgaacgagg gacaccctga caagctctgc gaccaaactc ccgatgctgt 180  
 octcgacgct tgccttgaac aggaccaga cagcaagggt gcctgcgaaa catgcaccaa 240  
 gaccaacttg gtcattggtct tcggagagat caccaccaag gccaacgttg actacgagaa 300  
 gatcgtgogt g 311

<210> 1920  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1920  
  
 ngcangcacg cgtacgtaag ctcggaattc ggctcgaggc ctgagcagg acccagacag 60  
 caaagttgcc tgcgaaacat gcacaaaac caacttggtc atggtcttcg gagaaatcac 120  
 gaccaaggcc aacgttgact acgagnagat agtgcgtgac acctgcagga acatcggtt 180  
 cgtctcaaat gatgtgggac tggatgccga caactgcaag gtctctgtca acattgagca 240

gcagancct gatattgctc aggggtgnnc ccggccacct t 281

<210> 1921  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1921

tccgaagtct cangcacgcy tacgtaagct cgggaattcgg ctcgagntac ctatgggtggg 60  
 tgggggtgctc atgggtggagg tgccttttca ggggaaggacc ctaccaaggt tgacagaagt 120  
 ggtgcctata tcgtgaggca ggctgcaaag agtggtgtgg caaatggcct tgcnaaaang 180  
 gtgcnntggc cnangttttn aaggccatng gtgtccctga gcccttgtca gtgtttgtgg 240  
 acacttatgg aactgggaag attcctgaca aggnngnttct tcaaattgtg aaggngantt 300  
 cngncttcng acntg 315

<210> 1922  
 <211> 259  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1922

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ggtggagggtg ccttttcagg 60  
 gaaggaccct accaaggttg acagaagtgg tgcctatata gtgaggcagg ctgcaaagag 120  
 tggtgtggca aatggccttg cagaagggtgc attgtccaag tttcctatgc cattgggtgtc 180  
 cctgagccct tgtcagtgtt tgnccgacact tatggaactg ggaagattcc tgacaaggag 240  
 attcttcaaa ttgtgaagg 259

<210> 1923  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1923

agtcgcatgc acgcgtacgt aagctcggga attcggctcg agacctctca agtttttgaa 60  
 gtatagagat ggcagngaca ttcctattta cctcagagtc agtgaacgag ggacaccctg 120  
 acaagctctg tgaccaaatc tcatgctgtc ctcgacgctt gcctcgaaca ggaccagac 180

agcaaggttg cctgcgaaac atgcaccaaa accaacttgg tcatggtctt cggagaaatc 240  
acgaccaagg ccaatgttga ctacgagaag atagtgcgtg acacctgcag gaacatcggt 300

<210> 1924  
<211> 290  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1924

anncgcangc acgcgtacgt aagctcggaa ttcggctcga gtcaagtttn tgaagtatag 60  
agatggcaga gacattccta tttaacctcag agtcagtga cagaggacac cctgacaagc 120  
tctgtgacca aatctctgat gctgtcctcg acgcttgctt cgaacaggac ccagacagca 180  
aggttgcttg cgaacatgc accaaaacca acttggtcca tggctcttcgg agaaatcacg 240  
accaaggcca atgttgacta cgagaagata gtgcgtgaca cctgcaggaa 290

<210> 1925  
<211> 294  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1925

ngtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gcggtcgcag cggtcgcagg 60  
gcaaattggc ttgccagaag gtgcattgtc caagtttctt atgccattgg tgtcnctgag 120  
cccttgctcag tgtttgtgga cacttatgga actgggaaga ttcctgacaa ggagattctt 180  
caaattgtga aggagaattt cgacttcaga cctggaatga tcaccattaa cttggacctt 240  
aagaggggtg gccataggtt cctcaagaca gctgcttatg gacactttgg aagg 294

<210> 1926  
<211> 473  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1926

anctttgtac gcgccccagg taccggtaaa ggaattccng gctcgaccca cgcgtaagcc 60  
cacgcgtccg tacggctgcg agaagacgac agaagggggc agcgcttgat ttgaggccag 120  
gcaagcccca ctcaaccacc acacctctcc tcgttcacgc tacccttttc tgctcttctt 180

ctacctttca agtttttaaaa gtataaagat ggcagagaca ttcctattta cctcagagtc 240  
 ggtgaacgag ggacaccctg acgagctctg cgaccaaadc tccgatgctg tcctcgacgc 300  
 ttgcctcgag caggacccag acagcaaagt tgccctgcgaa acatgcacca aaaccaactt 360  
 ggtcatggtc ttccggagaaa tcacgaccaa ggccaacgtt gactacgaag aagatagtcg 420  
 gtgacacctg cangaacatc ggcttcgtct caaatgatgt tggaactgga tgc 473

<210> 1927  
 <211> 490  
 <212> nucleic acid  
 <213> Glycine max

<400> 1927

Sequence 1

atttnacctt cccnggccn gttaaaggta aaganttccc gggncgaccc acgcgtccgc 60  
 ccacgcgtcc gtacggctgc gagaagacga cagaaggggg gcagcgcttg atttgaggcc 120  
 aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt tctgctcttc 180  
 ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt tacctcagag 240  
 tcggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgcctcgac 300  
 gcttgccctg agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaaccaac 360  
 ttggtcatgg tcttcggaga aatcacgacc aaggccaacg ttgactacga agaagatagt 420  
 gcgtgacacc tgcaggaaca tcggcttcgt ctcaaaatga tgtgggactg gatgccgaca 480  
 actgnnangg 490

<210> 1928  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max

<400> 1928

aanagtcgca ngcacgctta cgtnaagctc ggaattcggc tcgagaaagc gggntactgt 60  
 ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga agttaaaatg 120  
 gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180  
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag 240  
 gttgcctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga gatcacaacc 300

aaggccaacg tggactatgn

320

<210> 1929  
<211> 294  
<212> nucleic acid  
<213> Glycine max

<400> 1929

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ggagattctc aacattgtga 60  
aggaaaactt tgatttcagg cctgggatga tctccatcaa ccttgatctc aagaggggtg 120  
gaaataacag gtttttgaag actgctgcct atggacactt tggaagagaa gaccctgact 180  
tcacatggga agtgggtcaaa cccctcaagt gggagaaggc ctaagtaatt cattccactg 240  
ctctatgctg gaagtttttt gagcggtgcc cttataatat gtctaataatc catn 294

<210> 1930  
<211> 304  
<212> nucleic acid  
<213> Glycine max

<400> 1930

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag ggagattctc aacattgtga 60  
aggaaaactt tgatttcagg cctgggatga tctccatcaa ccttgatctc aagaggggtg 120  
gaaataacag gtttttgaag actgctgcct atggacactt tggaagagaa gaccctgact 180  
tcacatggga agtgggtcaaa cccctcaagt gggagaaggc ctaagtaatt cattccactg 240  
ctctatgctg gaagtttttt gagcggtgcc cttataatat gtctaataatc cataactttc 300  
cacg 304

<210> 1931  
<211> 321  
<212> nucleic acid  
<213> Glycine max

<400> 1931

cgcacatgcacg cgtacgtnag ctcggaattc ggctcgagct tctacctctc aagtttttga 60  
agtatagacn ncggcagaga cattccctat ttaccttcag agttcagtga acgagggaca 120  
cnctgacaag ctctgtgacc aaatctctga tgctgtcctc gacggttgcc tcgaacagga 180



cccagacagc naggttgcct gcgaaacatg caccaaaacc aacttgggtca tgggtcttcgg 240  
 agaaatcacg accaaggcca atgttgacta cgagaagata gtgcgtgaca cctgcaggaa 300  
 catcggcttt gtctcaaacg a 321

<210> 1932  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1932

cgcatgcacg cgtacgtaag ctcggaattc ggctcgagct tctacctctc aagtttttga 60  
 agtatagaga tggcagagac attcctatctt acctcagagt cagtgaacga gggacaccct 120  
 gacaagctct gtgaccaaatt ctctgatgct gtctctgacg cttgcctcga acaggaccca 180  
 gacagcaagg ttgcctgcga aacatgcacc aaaaccaact tgggtcatggc cttcggagaa 240  
 atcacgacca aggccaatgt tgactacgag aaganagtgc g 281

<210> 1933  
 <211> 292  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1933

natacatgca cgcgtacgta agctcggaat tcggctcgag ctctctgctc ttcttctacc 60  
 tttcaagttt ttaaagtatt aagatggcag agacattcct atttacctca gagtcagtga 120  
 acgagggaca ccttgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 180  
 ttgaacagga cccagacagc aaggttgcct gcgaaacatg caccaagacc aacttgggtca 240  
 tgggtcttcgg agagatcacc accaaggcca acgttgacta cgagaagatc gt 292

<210> 1934  
 <211> 266  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1934

ctctgcgncc aaatctccga tgctgtcctc gacgcttgcc ttgaacagga cccagacagc 60  
 aaggttgcct gcgaaacatg caccaagacc aacttgggtca tgggtcttcgg agagatcacc 120

accaaggcca acgttgacta cgagaagatc gtgcgtgaca cctgnaggaa catcggttc 180  
 gtctcaancg atgtgggact tgatgctgac aactgccaaag gtnctgnaa acattgaggn 240  
 nncagagccc tggatattgc ccaggg 266

<210> 1935  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max

<400> 1935

cgcnaangc gtacgtnagc tcggaattcg gctcgagncg gggtactgtc tgttcaagct 60  
 accatctctc tctctctttc ttagtgctc cttgccagaa gttaaaatgg cccaagaaac 120  
 tttcctattc acatctgaat cagtgaacga ggggcaccct gacaagctct gtgaccagat 180  
 ctccgatgct gtgctgatg catgcttga gcaggaccct gacagcaagg ttgcctgtga 240  
 aacctgcacc aagaccaaca tggatgatgg tttcggagag atcacaacca aggccaacgt 300  
 ggactatgag 310

<210> 1936  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max

<400> 1936

gtgcangca cgcgtacgta agctcggaat tcggctcgag gcaacagagt cctgatattg 60  
 ctcaagggtgt gcacggccac ctcaaaaga ggctgagga gattggtgct ggtgaccaag 120  
 gtcatatgtt cggctatgcc actgacgaga ctcccgagct catgcccttg agccatgtcc 180  
 ttgccacgaa gctcgggtgcc aagctcaccg aggttcggaa gaacgggaca tgcccttggc 240  
 tgagacctga tggaccactg ntgantgatt acgatcacga ttaattcggc cccgacagt 299

<210> 1937  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max

<400> 1937

ncnacgcan gcacgcgtac gtnagctcgg aattcggctc gagctctcaa gtttttgaag 60

tatagagatg gcagagacat tcctatTTtac ctcagagtca gtgaacgagg gacaccctga 120  
 caagcttctg tgaccnaaat ctctgatgct gtccctcgacg cttgcctcga acaggaccca 180  
 gacagcaagg ttgcctgcga aacatgcacc aaaaccaact tggatcatggt cttcggagaa 240  
 atcacgacca aggccaatgt tgactacgag aagatagtgc gtgacacctg caggaacatc 300  
 ggctttgtct t 311

<210> 1938  
 <211> 319  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1938

gatgcacgcg tacgtaagct cggaattcgg ctcgagcaca aagcgggtta ctgtctgttc 60  
 aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa aatggcccaa 120  
 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180  
 cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag caangttgcc 240  
 tgtgaaacct gcaccaagac caacatggta tggttttcgg agagatcaca accaaggcca 300  
 acgtggacta tgagaagat 319

<210> 1939  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1939

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cggctcgagg cacgctctgc 60  
 ttccagcgag tgttctttct tcgtttcaac accttaattt gcanacgctg cttcttcccg 120  
 cttgagaaat ggcacaagaa acctttctat tcacatctga atctgtaaac gagggtcacc 180  
 ccgacaagct gtgcgaccag atctctgatg cagtgcctga tgcgtgcott gaacaggacc 240  
 ctgacagcaa ggttgctgtg gagacatgca ccaagaccaa catggtcatt gtctttggag 300  
 agatcacaac caagg 315

<210> 1940  
 <211> 303  
 <212> nucleic acid

<213> Glycine max

<400> 1940

cgcangcacg cgtacgtaag ctcggaattc ggctcgaggt cttgatgctg acaactgcaa 60  
ggtcctcgtc aacattgagc aacagagtcc tgatattgct caaggtgtgc acggccacct 120  
caciaagagg cctgaggaga ttggtgctgn tgaccaaggt catatgttcg gctatgccan 180  
tganganact cccgagctca tgccttgag ccatgtcctt gccacgaagc tcggtgccaa 240  
gtctcaccga ggctnggnag aacgggacat ccctgggnt gagacntgnt ggcaaagncc 300  
aaa 303

<210> 1941

<211> 335

<212> nucleic acid

<213> Glycine max

<400> 1941

tcgctgcacg cgtacgtaag ctcggaattc ngctcgaggc ccactcaac caccacacct 60  
ctcctcgctt acgtacccc tttctgctct tttctacct ttcaagtttt aaaagtataa 120  
agatggcaga gacattccta tttacctcag agtcggtgaa cgaggacac cctgacaagc 180  
tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca 240  
aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaaatcacga 300  
ccaaggccaa cgttgactac gagaagatag tgcgt 335

<210> 1942

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 1942

tcgcangcnc gcgtacgtna gctcggaatt cggctcgagg ggattctcaa cattgtgaag 60  
gaaaactttg atttcaggcc tggtatgatc tccatcaacc ttgatctcaa gaggggtgga 120  
aataacaggt ttttgaagac tgctgcctat ggacactttg gaagagaaga ccctgacttc 180  
acatgggaag tggtcaaacc cctcaagtgg gagaaggcct aagtaattca ttccactgct 240  
ctatgctgga agttttttga gcgttgccct ataatatgtc taata 285

<210> 1943  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1943  
  
 ngctctcangc acncgtacgt aagctnggaa ttcggctcna ggtctgttcn ngctaccatc 60  
 tctctnctct ctttcttagt gctccttgcc cagaagttaa antggcccaa gaaactttcc 120  
 tattcacatc tgaatnagt aacgaggggc accctgacaa gctctgtgac cagatctccg 180  
 atgctgtgct cgatgcatgc ttggagcagg accctgacag canngttgcc tgtgaaacct 240  
 gcaccaagac caacatgggtg atggttttcg gagagatcac naccaaggnc aacgtggact 300  
 atgagaagat 310

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<210> 1944  
 <211> 317  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1944  
  
 gtgcgatgca cgcgtacgta agctcgggaa ttcggctcga ggtaggttc tgcacgctct 60  
 gcttccagcg agtggttcttt cttcgtttca acaccttaat ttgcacacgc tgctttctca 120  
 gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa cgaggggtcac 180  
 cccgacaagc tgtgcgacca gatctctgat gcagtgtcgc atgcgtgcct tgaacaggac 240  
 cctgacagca aggttgacctg tgagacatgc accaagacca acatgggtcat ggtcttttga 300  
 gagatcacia ccaaggc 317

<210> 1945  
 <211> 331  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1945  
  
 tngnnngcac gcgtacgtaa gctcgggaatt cggctcgagc ggctcgagtt tgggagttag 60  
 gttctgcacg ctctgcttcc agcgagtgtt ctttcttcgt ttcaacacct taatttgcac 120  
 acgctgcttc ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180

taaacgaggg tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctcgatgcgt 240  
gccttgaaca ggaccctgac agcaaggttg cctgtgagac atgcaccaag accaacaatgg 300  
tcattggtctt tggagagatc acaaccaagg c 331

<210> 1946  
<211> 314  
<212> nucleic acid  
<213> Glycine max

<400> 1946

nncacgcgta cgtaagctcg gaattcggct cgagnggagt taggttctgc acgctctgct 60  
tccagcgagt gttctttctt cgtttcaaca ccttaatttg cacacgctgc ttcttcngct 120  
tgagaaatgg cacaagaaac ctttctatct acatctgaat ctgtaaacga gggtcacccc 180  
gacaagctgt ggcaccagat ctctgatgca gtgctcgatg cgtgccttga acaggaccct 240  
gacagcaagg ttgcctgtga gacatgcacc aagaccaaca tggatcatggc ctttggagag 300  
atcacaacca aggc 314

<210> 1947  
<211> 306  
<212> nucleic acid  
<213> Glycine max

<400> 1947

ctcgnatgca cgcgtacgta agctcggaat tcggctcgag gcacgctctg cttccagcga 60  
gtgttctttc ttcgtttcaa caccctaatt tgcacacgct gcttcttcag cttgagaaat 120  
ggcacaagaa acctttctat tcacatctga atctgtaaac gagggtcacc ccgacaagct 180  
gtgcgaccag atctctgatg cagtgcctga tgcgtgcctt gaacaggacc ctgacagcaa 240  
ggttgcctgt gagacatgca ccaagaccaa catggatcatg gtctttggag agatcacaac 300  
caaggc 306

<210> 1948  
<211> 297  
<212> nucleic acid  
<213> Glycine max

<400> 1948

atngnagtcg cangcncgcg tacgt nagct cggaattcgg ctcgaggnga ttctcaacat 60  
 tgtgaaggaa aactttgatt tcaggcctgg tatgatctcc atcaaccttg atctcaagag 120  
 ggggtggaaat aacaggtttt tgaagactgc tgcctatgga cactttggaa gagaagaccc 180  
 tgacttcaca tgggaagtgg tcaaaccct caagtgggag aaggcctaag taattcattc 240  
 cactgctcta tgctggaagt tttttgagcg ttgcccttat aatatgtcta atatcca 297

<210> 1949  
 <211> 217  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1949

gcgtacgtaa gctcggaatt cggctcgagg acaaatgcaa ggtgttggtc aacattgagc 60  
 agcagagccc tgatatcgcc caggggtgtgc acggtcactt caccaagcgc ccagaggagg 120  
 ttggtgctgg tgaccaggtt cacatgtttg gctatgccac tgatgaaacc cctgagtaca 180  
 tgccctcag ccatgtcctt gcaaccaaac tcgggtgc 217

<210> 1950  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1950

agnnaangca cgcgtacgta agctcggaat tcggctcgag actctctctg ctcttcttct 60  
 acctttcaag tttttaaagt attaagatgg cagagacatt cctatttacc tcagagtcag 120  
 tgaacgaggg acaccctgac aagctctgcg accaaatctc cgatgctgtc ctgacgctt 180  
 gccttgaaca ggaccagac agcaagggtg cctgcgaaac atgcaccaag accaacttgg 240  
 tcatggtctt cggagagatc accaccaagg ccaacgttga ctacgagaag a 291

<210> 1951  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1951

gtcgcanntt ngcgtacgta agctcggaat tcggctcgag actctctctg ctcttcttct 60

acctttcaag tttttaaggt attaagatgg cagagacatt cctatattacc tcagagtcag 120  
tgaacgaggg acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt 180  
gccttgaaca ggaccagac agcaaggttg cctgcgaaac atgcaccaag accaacttgg 240  
tcattggtctt cggagagatc accaccaagg ccaacgttga ctacgagaag a 291

<210> 1952  
<211> 319  
<212> nucleic acid  
<213> Glycine max  
<400> 1952

gtgcgatgca cgcgtacgta agctcggaat tcggctcgag gttaggttct gcacgctctg 60  
cttcacgaga gtgttctttc ttcgtttcaa caccttaatt tgcacacgct gcttcttcag 120  
cttgagaaat ggcacaagaa acctttctat tcacatctga atctgtaaac gagggtcacc 180  
ccgacaagct gtgcgaccag atctctgatg cagtgtctga tgcgtgcctt gaacaggacc 240  
ctgacagcaa ggttgctgtg gagacatgca ccaagaccaa catggtcatg gtctttggag 300  
agatcacaac caaggccag 319

<210> 1953  
<211> 288  
<212> nucleic acid  
<213> Glycine max  
<400> 1953

gtcgnangca cgcgtacgta agctcggaat tcggctcgag tctctctgtt ctcttctacc 60  
tctcaagttt ttgaagtata gagatggcag agacattcct atttacctca gagtcagtga 120  
acgagggaca ccctgacaag ctctgtgacc aaatctctga tgctgtcctc gacgcttgcc 180  
tcgaacagga ccagacagc aaggttgctt gcgaaacatg caccaaaacc aacttggtca 240  
tggtcttcgg agaaatcacg accaaggcca atgttgacta cgagaaga 288

<210> 1954  
<211> 248  
<212> nucleic acid  
<213> Glycine max  
<400> 1954



acctctcaag tttttgaagt atagagatgg cagagacatt cctattttacc tcagagtcag 60  
tgaacgaggg acaccctgac aagctctgtg accaaatctc tgatgctgtc ctogacgctt 120  
gcctcgaaca ggaccagac agcaagggtg cctgcgaaac atgcaccaa accaacttgg 180  
tcatggtctt cggagaaatc acgaccaagg ccaatgttga ctacgagaag atatgctga 240  
cactgcag 248

<210> 1955  
<211> 309  
<212> nucleic acid  
<213> Glycine max  
<400> 1955

cgcangcacg cgtacgttag ctcggaattc ggctcgagnt ttgaaccctt ctggcaggtt 60  
tgtcattgga gggccgcatg gngatgctgg tctcaccggc ngcaagntca ncatcgacac 120  
ctatgntcng atgggggtgca catggtggtg gtgccttctn tgggaaggat ccgccaangt 180  
tgataggagt ggtgcctaca ttgtgaggca agctgcaaag agcattgttn caaatggant 240  
tgctaggagg gcaattgtgc aagtttcta tgccattggt gtgcctganc cntgtctgtg 300  
nttgtnac 309

<210> 1956  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
<400> 1956

cgtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gggaggaggt gccttttcag 60  
ggaaggaccc taccaaggtt gacagaagtg gtgcctatat cgtgaggcag gctgcaaaga 120  
gtgttgtggc aaatggcctt gccagaaggt gcattgtcca agtttctat gccattggtg 180  
tcctgagcc ctgtcagtgt ttgtggacac ttatggaact gggaagattc ctgacaagga 240  
gattcttcaa atgtgaagga gaattcgact tcagacctgg aatgatcacc at 292

<210> 1957  
<211> 317  
<212> nucleic acid  
<213> Glycine max

<400> 1957

tgcatgcacg cgtacgtaag ctcggaattc ggctcgagta acaacagcac aaagcgggtt 60  
actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg ccagaagtta 120  
aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctgaca 180  
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 240  
gcaagggttg ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca 300  
caaccaaggc caacgtg 317

<210> 1958

<211> 219

<212> nucleic acid

<213> Glycine max

<400> 1958

tagtgectcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60  
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
atgottggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
ggatgatggt ttcggagaga tcacaaccaa ggccaacgt 219

<210> 1959

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 1959

tcgatgcaen cgtacgtgag ctcggaattc ggctcgaggc gggtnnctgt ctgttcaagc 60  
taccatctct ctctctcttt cttagtgcct cnttgccaga agttaaagt gccaagaaa 120  
ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc tgtgaccaga 180  
tctccgatgc tgtgctcgat gcatgcttg agcaggaccc tgacagcaag gttgcctgtg 240  
aaacctgcac caagaccaac atggtgatgg ttttcggaga ggtcacaacc aaggccaacg 300  
tgatatgan aagattgtgc gtgac 325

<210> 1960

<211> 316

<212> nucleic acid

<213> Glycine max

<400> 1960

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcacaaagcg ggttactgtc 60  
tggtcaagct accatctctc tctctcttct ttagtgcttc cttgccagaa gttaaaatgg 120  
cccaagaaac ttctctattc acatctgaat cagtgaacga ggggcaccct gacaagctct 180  
gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240  
ttgcctgtga aacctgcacc aagaccaaca tggatgatgg ttctggagag atcacaacca 300  
aggccaacgt ggactn 316

<210> 1961

<211> 495

<212> nucleic acid

<213> Glycine max

<400> 1961

ggnnnnnnnn aatttactct gcccgncagg tacangtaca gaattcccgg ntcgaccac 60  
gcgtcagtac ggctgcgaga agacgacaga agggggcagc gcttgatttg aggccaggca 120  
agccccactc aaccaccaca cctctcctcg ttcacgctac ccctttctgc tcttcttcta 180  
cctttcaagt tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt 240  
gaacgaggga caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg 300  
cctcgagcag gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttgg 360  
catggtcttc ggagaaatca cgaccaangc caacgttgac tacganaaga tatgcgtgac 420  
acctgcaagg aacatcggtc tctctcaaat gatgttgga ctggatgccg acaactgcaa 480  
ggtctcgtca acatt 495

<210> 1962

<211> 270

<212> nucleic acid

<213> Glycine max

<400> 1962

agtcgcnaca cgcgtacgta aactcggaat tcggctcnag cnaggttgat aggnatgg 60  
cttacattgt gagacaggct gctaaganca ttgtggcaag tggacttgcc agaagggtgca 120

ttgtagcaag tgtcttange cattggtgtg cctgagcctt tgtctgtgtt tnttgacacc 180  
tatggcactg ggaagatcca tgataaggag attctcaaca ttgngaagga aaactttgat 240  
ttcangcctg gnatgatctc catcaacctt 270

<210> 1963  
<211> 282  
<212> nucleic acid  
<213> Glycine max  
<400> 1963

tgcaacaaaa ccaacttggg catggtcttc ggagaaatca cgancaaggc caacgnttga 60  
ctaaggagaa gnnatgcgnt gaacacctgg caggncctac ggcttcgtct caaatnangt 120  
gggactgccc tgccgacaac tgcaagggtc tcgtcaacat tgagcagcag agccctgata 180  
ttgctcaggg tgtacacggc caccttacca aaaaacctga agaaattggg gctgggtgacc 240  
agggtcacat gtttggcnat gccactgatg aaaccctga ct 282

<210> 1964  
<211> 306  
<212> nucleic acid  
<213> Glycine max  
<400> 1964

nngttcntgc acgcgtacgt cagctcggaa ttcggctcga ggcggttac tgtctgttca 60  
agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaa atggccaag 120  
aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180  
agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggtgcct 240  
gtgaaacctg caccaagacc aacatgggtga tggttttcgg agagtnhaca accaaggcca 300  
acgtgg 306

<210> 1965  
<211> 317  
<212> nucleic acid  
<213> Glycine max  
<400> 1965

gcangcacgc gtacgtaagc tcggaattcg gctcgaggac ttaacaacag cacaaagcgg 60

gttactgtct gttcaagcta ccactctctct ctctctttct tagtgctcc ttgccagaag 120  
 ttaaaatggc ccaagaaact ttcctattca catctgaatc agtgaacgag gggcaccttg 180  
 acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccttg 240  
 acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggtt ttcggagaga 300  
 ncacaaccaa ggccaag 317

<210> 1966  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max

<400> 1966

togtangtaa gctcggaatt cggctcgagg cgggncactg tctgttcaag ctaccatctc 60  
 tctctctctt tcttagtgcc tccttgccag aagttaaaat ggcccaagaa actttcctat 120  
 tcacatctga atcagtgaac gaggggcacc ctgacaagct ctgtgaccag atctccgacn 180  
 ctgtgctoga tgcattgctg gancaggacc ctgacagcaa ggttgctgtg gaaacctgca 240  
 ccaagaccaa catggtgatg gttttcggag agatcacaac caaggccaac gtggat 296

<210> 1967  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max

<400> 1967

acgtcgcatg ctagcgtacg taagctcgga attcggctcg agcacaaagc gggttactgt 60  
 ctgttcaagc taccatctct ctctctcttt cttagtgctt ccttgccaga agttaaagt 120  
 gccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180  
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag 240  
 gttgctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga gatcacaacc 300  
 aaggccacgt ggactatg 318

<210> 1968  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max

<400> 1968

gtcncatgca cgcgtacgta agctcggaat tcggtctgag acagcacaaa gcgggttact 60  
gtntgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120  
tgggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 180  
ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc 240  
aaggttgcct gtgaaacctg caccaagacc aacatggtga tggttttcgg agagatcaca 300  
accaaggcca acg 313

<210> 1969

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 1969

ncgtcgcatg cagcgtacg taagctcgga attcggctcg agngcagttt taaaagtata 60  
aagatggcag agacattcct atttacctac agagtcggtg aacgagggac accctgacaa 120  
gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 180  
caaagttgcc tgcgaaacat gcacaaaaac caacttggtc atggtcttcg gagaaatcac 240  
gaccaaggcc aacgttgact acgagaagat agtgcgtgac acctgcagga a 291

<210> 1970

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 1970

ntgcantnac gcgtacgtaa gctcggaatt cggctcnagn cagacttaac aacagcacia 60  
agcgggttac tgtctgttca agctaccatc tctctcncct ctttcttagt gcctccttgc 120  
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180  
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
accctgacag caaggttgcc tgtgaaacct gcancaagac caacatggtg atggttttcg 300  
gagagatcac aaccaaggcc aacgtgg 327

<210> 1971

<211> 294  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1971  
  
 tcgcatgcac gcgtacgtaa gctcggaatt cggtcgcagc cacaccactc tctctgnntc 60  
 ttctttctacc tttcaagtnt ttaaagtatt aagatggcag agacattcct atttacctca 120  
 gagtcagtga acgagggaca ccttgacaag ctctgcgacc aaatctccga tgctgtcctc 180  
 gacgcttgcc ttgaacagga ccagacagc aagggtgcct gcgaaacatg caccaagacc 240  
 aacttggtca tgggtcttcgg agagatcacc accaaggcca acgttgacta cgag 294

<210> 1972  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1972  
  
 gtgcgatgca gcgtacgta agctcggaat tcggctcgag ntttcaagtt tttaaagtat 60  
 taagatggca gagacattcc tatntacctc agagtcagtg aacgagggac accctgacaa 120  
 gctctgcgac caaatctccg atgctgtcct cgacgcttgc cttgaacagg acccagacag 180  
 caaggttgcc tgcgaaacat gcaccaagac caacttggtc atggtcttcg gagagatcac 240  
 caccaaggcc aacgttgnet acgagaagtc gtgcgtgaca ctgaggaaca tcg 293

<210> 1973  
 <211> 339  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1973  
  
 tcgngcacg cgtacgtaag ctcggaattc ggctcgcagc agccatttgg gagttagggtt 60  
 ctgcacgctc tgcttcacgc gagtgttctt tcttcgnttc aacaccttaa tttgcacacg 120  
 ctgcttcttc agcttgagaa atggcacaag aaacctttct attcacatct gaatctgtaa 180  
 acgaggggtca ccccgacaag ctgtgcgacc agatctctga tgcagtgctc gatgcgtgcc 240  
 ttgaacagga ccctggacag caaggttgcc tgtgagacat gcaccaagac caacatggtc 300  
 atggtctttg gagagatcac aaccaaggcc aacgtagat 339

<400> 1974

[illegible]

<400> 1975

[illegible]

<400> 1976



tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc 240  
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga 300  
 gatcacaacc aaggc 315

<210> 1977  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max

<400> 1977

gtgcgancga cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcacaaag 60  
 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120  
 aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180  
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctt gagcaggacc 240  
 ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300  
 agatcacaac caaggc 316

<210> 1978  
 <211> 309  
 <212> nucleic acid  
 <213> Glycine max

<400> 1978

nnagangcac gnacaacgta agctcggaat tcggctcgag caacagcaca aagcgggtta 60  
 ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa 120  
 aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa 180  
 gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240  
 caaggttgcc tgtgaaacct gcaccaagac caacatgggt atggttttcg gagagatcac 300  
 aaccaaggc 309

<210> 1979  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max

<400> 1979

natgnntacg tnagctcgga attcggctcg agcagcacia agcgggttac tgtctgttca 60  
 agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaaa atggcccaag 120  
 aaacttttct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180  
 agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggtgcct 240  
 gtgaaacctg caccaagacc aacatggtga tggttttcgg agagatcaca accaaggc 298

<210> 1980  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1980

togcangcac gcgtacgtaa nctcggaatt cggtctgagn ttgggagtta ggttctgcac 60  
 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120  
 cttcagcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180  
 gtcaccccga caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccttgaac 240  
 aggacctga cagcaagggt gctgtgaga catgcacaa gaccaacatg gtcattggtct 300  
 ttggagagat caca 314

<210> 1981  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 1981

gtgcgatgca cgcgtacgta agctcggaat tcgggctcga gcttaacaac agcaciaaagc 60  
 gggttactgt ctgttcaagc taccatntct ctctctcttt cttagtgcct ccttgccaga 120  
 agttaaaatg gcccaagnaa acttttctat tcacatctga atcagtgaac gaggggcacc 180  
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc 240  
 ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300  
 agatcacaac caaggccaac gtggn 325

<210> 1982  
 <211> 315  
 <212> nucleic acid

<213> Glycine max

<400> 1982

gtcgcangca cgcgtacgta agctcggaat tcggctcgag nttgggagtt aggtttctgca 60  
cgctctgctt ccagcgagtg ttctttcttc gtttcaacac cttaatttgc acacgctgct 120  
tcttcagctt gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag 180  
ggtcaccccg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa 240  
caggaccctg acagcaaggt tgctgtgag acatgcacca agaccaacat ggtcatggtc 300  
tttgagagaga tcaca 315

<210> 1983

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 1983

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cctcagccat gtccttgcaa 60  
ccaaacttgg tgctcgcttc acagagggtta ggaagaatgg cacctgtgct tggttgaggc 120  
cagatggtaa gacacaagta accgtcgagt actacaatga caatgggtgcc atggttccag 180  
ttcgtgtcca cactgtccta atttccaccc aacatgatgn ncctgtgagc aatgatcaaa 240  
ttgtgcgga cttaaaggca tgttataaac ctgncatccn ggaaaatact tgnaggaa 298

<210> 1984

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 1984

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcgggttact gtctgttcaa 60  
gtaccatct ctctctctct nnccttagtg ctccttgcc agaagttaa atggccaag 120  
aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180  
agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aagggtgcct 240  
gtgaaacctg caccaagacc aacatgggtga tggttttcgg agagatcaca accaaggcc 299

<210> 1985

<211> 306  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1985  
  
 ncgtnncntn nagctcggaa ttcggctcga gcttaacaac agcacaaagc gggttactgt 60  
 ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga agttaaaatg 120  
 gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180  
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag 240  
 gttgcctgtg aaacctgcac caagaccaac atggtgatgg ttttcggaga gatcacaacc 300  
 aaggcc 306

<210> 1986  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1986  
  
 gtencatgca cgcgtacgta agctcggnat tcggctcgag aagcgggtta ctgtctgttc 60  
 aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa aatggcccaa 120  
 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180  
 cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag caaggttgcc 240  
 tgtgaaacct gcaccaagac caacatggtg atggttttcg gagagatcac aaccaaggcc 300

<210> 1987  
 <211> 319  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 1987  
  
 gaaccgtngc tggtagctaa gctcggaatt cggctcgagg cagacttaac aacagcacia 60  
 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120  
 agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240  
 ccctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300

agagatcaca accaaggcc 319

<210> 1988  
<211> 311  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1988

gnannngcac gcgtacgtaa gctcggaatt cggctcgaga acaacagcac aaagcgggtt 60  
actgtctgtt caagctacca tctctctctc tctttcttag tgctctcttg ccagaagtta 120  
aatgggcca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctgaca 180  
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 240  
gcaaggttgc ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca 300  
caaccaaggc c 311

<210> 1989  
<211> 331  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1989

ctngcangca gcgtacgtaa gctcggaatt cggctcgagg cagacttaac aacagcacia 60  
agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120  
agaagttaaa atgggcccac ganactttcc tntcacatc tgaatcagt aacgaggggc 180  
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttcg 300  
gagagatcac aaccaaggcc aacgtggact a 331

<210> 1990  
<211> 319  
<212> nucleic acid  
<213> Glycine max  
  
<400> 1990

tcaacagtcg catgcacgcg tacgtaagct cggaattcgg ctcgagaaca acagcaciaa 60  
gcgggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctccttgcca 120

gaagttaaaa tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac 180  
cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac 240  
cctgacagca aggttgcttg tgaaacctgc accaagacca acatgggtgat ggttttcgga 300  
gagatcacia ccaaggccc 319

<210> 1991  
<211> 288  
<212> nucleic acid  
<213> Glycine max  
<400> 1991

ntaangcacg cgtacgtaag ctcggaattc ggctcgaggg acaaatgcaa ggtgttggtc 60  
aacattgagc aacagagccc ggatatcgcc cagggtgtgc acggccactt caccaagcgc 120  
ccagaggagg ttggtgctgg tgaccagggt cacatgtcac angtatgcc ncatgncac 180  
ccccgagtac atgcccctca gccatgtcct tgcaacaaaa cttggtggnt cgcncacag 240  
aggttaggag aattgcactg tgcttggttg aggcagatg gtaagaca 288

<210> 1992  
<211> 333  
<212> nucleic acid  
<213> Glycine max  
<400> 1992

nnncnngcac gcgtacgtng ctcggaattc ggctcgagct taacaacagc acaaagcggg 60  
ttactgtctg ttcaagctac catctctctc tctctttctt agtgcctcct tgccagaagt 120  
taaaatggcc caagaaactt tcctattcac atctgaatca gtgaacgagg ggcaccctga 180  
caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga 240  
cagcaagggt gctgtgaaa cctggcacca agaccaacat ggtgatggtt ttcggagaga 300  
tcacaaccaa ggccaagtgg actatgagaa gat 333

<210> 1993  
<211> 325  
<212> nucleic acid  
<213> Glycine max  
<400> 1993

tcgcatgcac gcgtacgtaa gctcgggaatt cggctcgagg ttgagaccaa gacacactcg 60  
 ttcatatatc tctctgtctt tctcttctct tctacctctc aagtttttga agtataaaga 120  
 tggcagagac attcctattc acctcggagt cagtgaacga gggacaccct gataagctct 180  
 gcgaccaatc tccgatgctg tctcgcacgc ttgcctcgaa caggacccag acagcaaggt 240  
 tgccctgcgaa acatgcacca agaccaactt ggtcatgggc ttccggagaga tcaccaccaa 300  
 ggccaacgtt gactacgaga agatc 325

<210> 1994  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 1994

acgtcgcacg cagcgcgtacg taagctcggg attcggctcg aggttactgt ctgttcaagc 60  
 taccatctct ctctctcttt cttagtgcct ccttgccaga agtaaaaatg gcccaagaaa 120  
 ctttctctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc tgtgaccaga 180  
 tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tgacagcaag gttgcctgtg 240  
 aaacctggca ccaagaccaa catggtgatg gttttcggng agatcacaac caaggccaag 300

<210> 1995  
 <211> 322  
 <212> nucleic acid  
 <213> Glycine max

<400> 1995

gcgtacgtaa gctcgggaatt cggctcgagn aagcnccact tcaaccacca cacnactctc 60  
 tctgctcttc ttctaccttt caagttttta aagtattaag atggcagaga cattcctatt 120  
 tacctcagag tcagtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc 180  
 tgtcctcgac gcttgcttgg aacaggaccc agacagcaag gttgcctgcg aaacatgcac 240  
 caagaccaat tggatcatgg cttcggagag atcaccacca aggccaagtt gactacgaga 300  
 agatcgtgcg tgacactgca gg 322

<210> 1996  
 <211> 321  
 <212> nucleic acid

<213> Glycine max

<400> 1996

tgggagtttag gttctgcacg ctctgcttcc agcgagtgtt ctttcttctgt ttcaacacct 60  
taatttgcac acgctgcttc ttcagcttga gaaatggcac aagaaacctt tctattcaca 120  
tctgaatctg taaacgaggg tcaccccgac aagcgtgcga ccagatctct gatgcagtgc 180  
tcgatgcgtg ccttgaacag gacctgaca gcaagggtgc ctgtgagaca tgcaccaaga 240  
ccaacatggt catggtcttt ggagagatca caaccaaggc aacgtagata tgagaagatg 300  
tcgtgnacat gcgcgaattg g 321

<210> 1997

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 1997

tcgcangcac gcgtacgtaa gctcgggaat tcggctcgan ctcgagccga ttcggctcga 60  
ggccttttca gggaaggacc ctaccacngg ttgacagaag tgaccgccta tattgtaagg 120  
cagncgtcaa agagtgttcg tgggcaaattg gccttgntag aagggtgcatt gtgcaagttt 180  
cctatgccat tgggtgtccct gagcccttgt cagtgtttgt ggacncttat ggaactggga 240  
agattcctga caaggagatt ctgcaaattg tgaaggagaa tttcgacttc agacctggna 300  
tga 303

<210> 1998

<211> 328

<212> nucleic acid

<213> Glycine max

<400> 1998

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ncagacttaa caacagcaca 60  
aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120  
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180  
acctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
acctgacag caagggtgcc tgtgaaacct gcaccaagac caacatggtg atgggtttctg 300



gagagatcac naccaaggcc aacgtggg

328

<210> 1999

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 1999

tcgcacacnt acgcncggaa tctcggcncg anaacagcac aaagcggggtt actgtctgtt 60

caagctacca tctctctctc tctntcttag tgctccctt gccagaagtt aaaatggccc 120

aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac aagctctntg 180

accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac agcaangttg 240

cctgtgaaac ctgcaccaag accaactgg tgatngtttt cggagagatc acaaccaagg 300

cncgg 305

<210> 2000

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2000

gtgcangca cgcgtacgta agctcggaat tcggctcgag cttacaaca gcacaaagcg 60

ggttactgtc tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa 120

gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180

gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240

gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300

atcacaacca ggccaagtgg a 321

<210> 2001

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2001

gtcgcatgca cgcgtacgta agctcgggaa ttcggctcga ggtgatttgg gagtttggag 60

cgactgaact aatcattaat ttgcactcgc tgtttcagct tcatcaccct tcttttgcac 120

catttatatc tcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180  
cgaggggtcac cccgacangc tgttcnanca gatctctgat gcagtacttg atgcgtgcct 240  
tgaacaggac cctgacagca aggttgcttg tgagacatgc accaagacca acatgggtcat 300  
ggtcttcgga gagatcacia ccaaggc 327

<210> 2002  
<211> 316  
<212> nucleic acid  
<213> Glycine max  
<400> 2002

ntcgnatncl agcltangtn agnnttcggn tclgagatttg ggagttaggt tctgcacgct 60  
ctgcttccag cgagtgttct ttcttcgttt caacacctta atttgcacac gctgctttct 120  
tengcttgag aaatggcnaca agnnaccttt ctattcacat ctgaatctgt aaacgaggggt 180  
caccocgaca anctgtgcga ccagatctct gatgcagtgc tclgatgcgtg ccttgaacag 240  
gacnctgaca gcaaggttgc ctgtgagaca tgcaccaaga ccaacatgggt catgggtcttt 300  
gganagatca caacca 316

<210> 2003  
<211> 334  
<212> nucleic acid  
<213> Glycine max  
<400> 2003

ccnclngccn acccltngnc nncntntcng tclgnngnnnc gtacgtnagc tclgnattclg 60  
gctclngcca agccccactc aaccaccaca ccactctctc tgctcttctt ctacctttca 120  
ngttttttaa gtattaagat ggcagagaca ttctattta cctcagagtc agtgaacgag 180  
ggacaccctg acaagctctg cgaccaaalc tccgatgctg tclctcgacgc ttgccttgaa 240  
caggaccclg acagcaaggt tgcltgclga acatgcacca agaccaactt ggtclatggtc 300  
ttcggagaga tcaccaccaa ggccaacgtt gact 334

<210> 2004  
<211> 216  
<212> nucleic acid  
<213> Glycine max

<400> 2004

tagtgctcc ttgccagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 60  
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
atgcttggag caggaccctg acagcaaggt tgctgtgaa acctggcacc aagaccaaca 180  
tggtgatggg tttcggagag atcacaacca aggcca 216

<210> 2005

<211> 319

<212> nucleic acid

<213> Glycine max

<400> 2005

gnacgacgca ngcacgcgta cgtnagctcg gaattcggct cgagctcatg gtgatgctgg 60  
tctcactgga agaaagatca tcattgatac ctatggtggg tggggtgctc atggtggagg 120  
tgcccttttca gggaaggacc ctaccaaggt tgncagaagt ggtgcctata tcgtgaggca 180  
ggctgcaaag agtggttngtg gcaaattggc ttgccagaag gtgcattgtc caagtttcct 240  
atgccattgg gtgtccctga gccctngtca gnggtnggtg gacattatgg nncntgggaa 300  
nttcctcaca aggggtttt 319

<210> 2006

<211> 295

<212> nucleic acid

<213> Glycine max

<400> 2006

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc tcgagccgct cgagccgatt 60  
cggctcgagg tggccctcat ggtgntgctg gtctcactgg accgaaagat acntcattga 120  
tacctatggg ggggtggggac ctcatgggtg aggtgccttt tcaggggaagg accctaccaa 180  
ggttgacaga agtggtgcct atatngtgag gcaggctgca aanagtgttg tggcaaattg 240  
ccttgccaga aggtgcattg tccaagtttc ctatgcnatt ggtgtccctg agccc 295

<210> 2007

<211> 261

<212> nucleic acid

<213> Glycine max

<400> 2007

tatctctctg ttctcttcta cctctcaagt ttttgaagta tagagatggc agagacattc 60  
ctatttacct cagagtcagt gaacgaggga caccctgaca agctctgtga ccaaattctct 120  
gatgctgtcc tcgacgcttg cctcgaacag gaccagaca gcaagggttg ctgcgaaaca 180  
tgcacaaaaa ccaacttggc catggtcttc ggagaatcac gaccaaggcc aatgtngant 240  
acgagaagat atgcgtgacc c 261

<210> 2008

<211> 422

<212> nucleic acid

<213> Glycine max

<400> 2008

caggcaagcn ccaactcaacc accacacctc tctngttca cgctacccgc tttctgctct 60  
tcttctacct ttcaagtttt aaaagtataa agatggcaga gacattccta ttacctcag 120  
agtcggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctng 180  
acgcttgctt cgagcaggac ccanacagca aagttgcctg cgaaacatgc accaaaacca 240  
acttggtcat ggtcttcgga gaaatcacga ccaaggccaa cgttgactac gaagaagata 300  
gtgcgtgaca cctgcaggaa ccattngnnt tngtctnaaa tgatgtgggg actggatgcc 360  
cgacaactgg caaggtcctc gtcnaacatt gancatcaaa agccctggtn ttggttnagg 420  
gg 422

<210> 2009

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 2009

tcgngcacn cgtacgtnag ctcggnnttc ggctcgacct cgagccgaat cggctcgagg 60  
ggttactgtc tgttcaagct aaccatctct ctctctctac tontagtgcc tcttgccan 120  
aagttaaaat ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180  
ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc 240  
ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag 300

agatcacaa

309

<210> 2010  
<211> 280  
<212> nucleic acid  
<213> Glycine max

<400> 2010

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<210> 2011  
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<400> 2011

atgcacgcgt acgtaagctc ggaattcggc tcgaggcaga cttacaaca gcacaaagcg 60  
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<211> 290  
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<400> 2012

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gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca aggttgcttg 240  
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<210> 2013  
 <211> 274  
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 <400> 2013

agtcgcannc acgcgtacgt aagctcggaa ttcggctcga nggctcgagc ggctcgnnggc 60  
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 gctatgccac tgacgagact nccgagctca tnncttgag cnatgtcctt gccacnaagc 180  
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<210> 2014  
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 gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240  
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 <213> Glycine max  
 <400> 2015

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gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240  
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 aaccaaggc 309

<210> 2016  
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 agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gaccctgaca 240  
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<210> 2017  
 <211> 294  
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 <400> 2017

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 gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180  
 cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag caaggttgcc 240  
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<210> 2018  
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 <400> 2018

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gccgcaagtt aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg 180  
gcaccctgac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttgagca 240  
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cggagagatc acaaccaagg c 321

<210> 2019  
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ttacatgtga 310

<210> 2021  
<211> 326  
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<400> 2021

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 actacgagaa atagtgcgtg acacct 326

<210> 2022

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2022

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 aatgacaatg gtgccatggt tccagttcgt gtccacactg tccataattc caccacaacat 180  
 gatgagnctg tgagcaatga tcaaagtctg cggaccttaa agagcatggt atcaagcctg 240  
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<210> 2023

<211> 545

<212> nucleic acid

<213> Glycine max

<400> 2023

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 nacattccta ttacctcaa gtagctgta acgagggaca ccctgacaag ctctgcgacc 240  
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 aacgttgact angagaaaga tngtgcgtga cacctgcagg aatatcggt tcgtctcagg 420

angatntggg acttgatnct gacatctgca angtecttgt aaacattncg cagcatancc 480  
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<210> 2024  
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 ctgacaagct ctgcgaccaa actccgatgc tgtctctgac gcttgcccttg aacaggaccc 180  
 agacagcaag gttgcctgcg aaacatgcac caagaccaac ttggtcatgg tcttcggaga 240  
 gatcaccacc aaggccaacg ttgactacga g 271

<210> 2025  
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 <212> nucleic acid  
 <213> Glycine max  
 <400> 2025

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 cgggtgaacga gggacaccct gacaagctct gcgaccaa atccgatgct gtcctcgacg 180  
 cttgcctcga gcaggaccca gacagcaaag ttgcctgcga aacatgnacc aaaaccaact 240  
 tggtcatggt cttcggagaa atcacgacca aggccaagtt gactacgaga agatagt 297

<210> 2026  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2026

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gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
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atcacaacca 310

<210> 2027  
<211> 310  
<212> nucleic acid  
<213> Glycine max  
<400> 2027

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gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300  
atcacaacca 310

<210> 2028  
<211> 309  
<212> nucleic acid  
<213> Glycine max  
<400> 2028

nngnancctta gagtcgcatg cagcgtacg taagctcgga attcggctcg aggttagggt 60  
ctgcacgctc tgcctccagc gagtggtctt tcttcgtttc aacaccttaa tttgcanacg 120  
ctgcttcttc ngcttgagaa atggcacaag aaacctttct attcacatct gaatctgtaa 180  
acgaggggtca ccccgacaag ctgtgcgacc agatctctga tgcagtgtc gatgcgtgcc 240  
ttgaacagga ccttgacagc aaggttgctt gtgagacatg caccaagacc aacatgggtca 300  
tggtctttg 309

<210> 2029  
<211> 487  
<212> nucleic acid  
<213> Glycine max

<400> 2029

aactctactt ggncaggccc cggtnacagaa atcccggctc gacccacgcg tcagtacggc 60  
tgcgagaaga cgacagaagg gggcagcgct tgatttgagg ccaggcaagc cccactcaac 120  
caccacacct ctctcgttc acgtacccc tttctgtctt tttctacct ttcaagtttt 180  
aaaagtataa agatggcaga gacattccta tttacctcag agtcgggtgaa cgagggacac 240  
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcangac 300  
ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 360  
gaaatcacga acaagggcaa cgttgactac gaaaaagata attgcntgac aacctgcagg 420  
gaacatcggc ttcgtctcaa atgatgttgg gactggatgc cgacaactgc aaaggtctcc 480  
gtcaaca 487

<210> 2030

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2030

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gccaaagccc actcaaccac 60  
cacaccactc tctctgtctt tttctacct ttcaagtttt taaagtatta agatggcaga 120  
gacattccta tttacctcag agtcagttaa cgagggacac cctgacaagc tctgcgacca 180  
aatctccgat gctgtcctcg acgcttgctt tgaacaggac ccagacagca aggttgcttg 240  
cgaaacatgc accaagacca acttggtcat ggtcttcgga gagatcacca ccaaggcc 298

<210> 2031

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2031

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ctgcttcacg cgagtgttct nacttcgttt caacacctta attgcacac gctgcttctt 120  
cagcttgaga aatggcacia gaaacctttc tattcacatc tgaatctgta aacgagggtc 180  
accccgacaa gctgtgcgac cagatctctg atgcagtgtc cgatgcgtgc cttgaacagg 240

accctgacag caaggttgcc tgtgagacat gcaccaagac caacatggtc atggtctttg 300  
g 301

<210> 2032  
<211> 297  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2032

cgtcgcangc acgcgtacgt nagctcggna ttccggctcga gngcacaaag cgggttactg 60  
tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat 120  
ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagct 180  
ctgtgaccag atctccgatg ctgtgctcga tgcattgctt gagcaggacc ctgacagcaa 240  
ggttgcctgt gaaacctgca ccaagaccaa catggtgatg gttttcggng agatcac 297

<210> 2033  
<211> 332  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2033

tgcacgcgta cgtaagctcg gaattcggct cgagatttga ggcaggcaa gcccnactca 60  
accaccacac ntctctctgt tnanctacc ctttctncc tcttcttcta cttttcaagt 120  
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 180  
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgctng cctcgagcag 240  
gaccagaca gcaaagttgc ctgcgaaaca tgcacaaaa ccaacttggc catggtcttc 300  
ggaganatca cgaccaaggc caacgttgac ta 332

<210> 2034  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2034

tcgcatgcac gcgtacgtna gctcggatt cggtctgagn acagcacaaa gcgggttact 60  
gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120

tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180  
tctgtgacca gatctccgat gctgtgntcg atgcatgctt ggagcaggac cctgacagca 240  
aggttgcttg tgaaacctgc accaagacca acatgggtgat ggtttttcgga gagatcacia 300

<210> 2035  
<211> 307  
<212> nucleic acid  
<213> Glycine max  
<400> 2035

ncgcangcac gcatacgtna gctcggaatt cggctcgagc tnaacaacag caciaaagcgg 60  
gttactgtct gttcaagcta ccactctctc tctctctttc ttagtgcttc cttgccagaa 120  
gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
gacagcaagg ttgctgtna aacctgcacc aagaccaaca tgggtgatgg tttcggagag 300  
atcacia 307

<210> 2036  
<211> 262  
<212> nucleic acid  
<213> Glycine max  
<400> 2036

ccaagcccca ctcaaccacc acaccactct ctctgctctt cttctacctt tcaagttttt 60  
aaagtattaa gatggcagag acattcctat ttacctcaga gtcagtgaac gagggacacc 120  
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctt gaacaggacc 180  
cagacagcaa ggttgcttg gaaacatgca ccaagaccaa cttggtcatg gtcttcggag 240  
agatcaccac caaggccaac gt 262

<210> 2037  
<211> 323  
<212> nucleic acid  
<213> Glycine max  
<400> 2037

aaatntanan gtcgcangca cgcgtacgta agctcggaat tcggctcgag cttacaaca 60

gcacaaagcg ggttactgtc tgttcaagct accatctcct ctctctcttt cttagtgcct 120  
 ccttgccaga agttaaagt gccaagaaa ctttcttatt cacatctgaa tcagtgaacg 180  
 aggggcaccc tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg 240  
 agcaggaccc tgacagcaag gttgcctgtg aaacctgcac caagaccaac atggtgatgg 300  
 ttttcggaga gatcacaacc aag 323

<210> 2038  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max

<400> 2038

gagagatcac a

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60  
 aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120  
 cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180  
 accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
 accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttgc 300  
 gagagatcac a 311

<210> 2039  
 <211> 301  
 <212> nucleic acid  
 <213> Glycine max

<400> 2039

ttcangcacn cgtacgtaag ctcggaattc ggctcgagca caaagcgggt tactgtctgt 60  
 tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt aaaatggccc 120  
 aagaaacttt cctattcaca tctgaatcag tgaacgagg gcacctgac aagctctgtg 180  
 accagatctc cgatgctgtg ctgatgcat gcttgagca ggacctgac agcaaggntg 240  
 cctgtgaaac ctgcaccaag accaactgg tgatggtttt cggngagatc acaaccaagg 300  
 n 301

<210> 2040  
 <211> 307  
 <212> nucleic acid

<213> Glycine max

<400> 2040

gtngcangca cgcgtacgta agctcggaat tcggctcgag cagcacaaag cgggttactg 60  
tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat 120  
ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagcc 180  
ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc ctgacagcaa 240  
ggttgctgtg gaaacctgca ccaagaccaa catggtgatg gttttcggag agatcacaac 300  
caaggcc 307

<210> 2041

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2041

cgcattgcagt ntacgtaagc tcggaattcn gctcgagcag cacaaagcgg gttactgtct 60  
gttcaagcta ccattctctc ctctctttct tagtgctcc ttgccagaag ttaaaatggc 120  
tcaagaaact ttctatttca catctgaatc agtgaacgag gaccacctg acaagctctg 180  
tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggacctg acagcaaggt 240  
tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttcggagaga tcacaacca 300  
ggc 303

<210> 2042

<211> 486

<212> nucleic acid

<213> Glycine max

<400> 2042

tngcnaactc ttacgcggtt caggtaccgg ttgnngaatt cccggggtcg acccacgcgt 60  
caagtacggc tgcgagaaga cgacagaagg gggcagcgt tgatttgagg ccaggcaagc 120  
ccactcaac caccacacct ctctcgttc acgtacccc tttctgctct tttctacct 180  
ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa 240  
cgaggggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgct 300



cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat 360  
 ggtcttcgga gaaatcacga ccaaggccaa cgttgactac gaagaagata gtgcgtnaca 420  
 cctgcagga acatccgnt nntnccaaaa tnangttga ncgggatccn anaatttgc 480  
 aggggt 486

<210> 2043  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2043

ngtcgcatgc acgcgtacgt aagctcgga ttcggctcga ggggcaccct gnacaagctc 60  
 tgtgaccaga tctccgatgc tgtgctcgat ggcattgctg gagcaggacc ctgacagcaa 120  
 ggttgctgt gaaacctgca ccaagaccaa catggtgatg gttttcggag agatcacaac 180  
 caaggccaac gtggactatg agcaagcttg tgnctgaca catgcaggaa cattggtttt 240  
 gtctctnatg atgtnggtct tggatgcna caactgcaag tctcgtcaac atngagcaac 300  
 agan 304

<210> 2044  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2044

gtcgcangca cgcgtacgta agctcgaatt cggctcgagg cagacttaac aacagcacia 60  
 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120  
 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240  
 ccctgacagc aaggttgctt gtgaacctgc accaagacca acatggtgat ggttttcgga 300  
 gagatcacia ccaggccang tggan 325

<210> 2045  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max

<400> 2045

gtcgcacatgca cgcgtacgta agctcggaat tcggctcgag tgagaaatgg cacaagaaac 60  
ctttctattc acatctgaat ctgtaaacga gggtcacccc gacaagctgt gcgaccagat 120  
ctctgatgca gtgctcgatg cgtgccttga acaggaccct gacagcaagg ttgcctgtga 180  
gacatggcac caagaccaac atggatcatg ttctttggag agatncacaa ccaagggcca 240  
acgtagacta tgagaagatt gttcctgnac acatgccggc gaantggatt ncannccg 298

<210> 2046

<211> 318

<212> nucleic acid

<213> Glycine max

<400> 2046

gtcgcacatgca cncgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60  
aagcgggtta ctgtctgttc aagctaccat ctctctctct cttctcttagt gcctccttgc 120  
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180  
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
accctgacag caagggttgc tgtgaaacct gcaccaagac caacatggtg aggttttcgg 300  
agagatcaca accaaggc 318

<210> 2047

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2047

gngtcgnang cagcgtacg tnagctcgga atgcggctcg aggggttact gtctgttcaa 60  
gctaccatct ctncctctct ttcttagtgc ctcttgcca gaagnnaaan tngcccaaga 120  
aactttccta ttcnnatctg aatcagtga cgaggggcac cctgacaagc tctgtgacca 180  
gatctccgat gctgtgctcg atgcatgcnt ggngcaggac nctgacagca aggttnoctg 240  
tgaaacntgc accaagacca acatggtgat ggttttcgga gagatcacia ccaaggccaa 300  
cg 302

<210> 2048

<211> 301  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2048  
  
 tcgnangcac gcgtacgtaa gctcggaatt cggctcnagt ttgggagtta ggttctgcac 60  
 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120  
 cttongcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180  
 gtcaccccgca caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccttgaac 240  
 aggaccctga cagcaagggt gctgtgaga catgcaccaa gaccaacatg gtcatgggtct 300  
 t 301

<210> 2049  
 <211> 273  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2049  
  
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 tctacctctc aagtttttga agtataaaga tggcagagac attcctattc acctcgaggt 120  
 cagtgaacga gggacaccct gataagctct gcgaccaa atccgatgct gtcctcgacg 180  
 cttgctcga acaggaccca gacagcaagg ttgctgcga aacatgcacc aagaccaact 240  
 tggatcatggt cttcggagag atcaccacca agg 273

<210> 2050  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2050  
  
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 agtgttcttt cttcgtttca acaccttaat ttgcanacgc tgcttcttct ggcttgagaa 120  
 atggcacaag aaacctttct attcacatct gaatctgtaa acgaggggtca ccccgacaag 180  
 ctgtgcgacc agatctctga tgcagtgtc gatgcgtgcc ttgaacagga ccttgacagc 240  
 aaggttgcct gtgagacatg caccaagacc aacatgggtca tggctcttga gagatcacao 300

ccagggccaa cgt

313

<210> 2051  
<211> 312  
<212> nucleic acid  
<213> Glycine max

<400> 2051

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcacaaag 60  
cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120  
aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180  
ctgacaanct ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc 240  
ctgacagcaa ggttgccctgt gaaacctgca ccaagaccaa catggtgatg gttttcggag 300  
agatcacaa ca 312

<210> 2052  
<211> 308  
<212> nucleic acid  
<213> Glycine max

<400> 2052

gcgtacgtaa gctcggaatt cngctcgagg cccactcaa ccaccacacc tctcctcggt 60  
cacgctaccc ctttctgctc ttcttctacc tttcaagttt taaaagtata aagatggcag 120  
agacattcct atttacctca gagtcggtga acgagggaca ccctgacaag ctctgcgacc 180  
aaatctccga tgctgtcctc gacgcttgnc tcgagcagga ccagacagc aaagttgcct 240  
gcgaaacatg caccaaaacc aacttggtca tggctctcgg agaaatcacg accaaggcca 300  
acgttgat 308

<210> 2053  
<211> 298  
<212> nucleic acid  
<213> Glycine max

<400> 2053

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ttccagcgag tggtctttct tcgtttcaac accttaattt gcacacgctg cttcttcagc 120

ttgagaaatg gcacaagaaa cctttctatt cacatctgaa tctgtaaacg agggtcaccc 180  
cgacaagctg tgcgaccaga tctctgatgc agtgctcgat gcgtgccttg aacaggaccc 240  
tgacagcaag gttgcctgtg agacatgnac caagaccaac atggatcatgg tctttggn 298

<210> 2054  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
<400> 2054

nanangangt cgcangcacg cgtacgtgag ctccgnattc ggctcgaggn aagccccact 60  
caaccaccac accactctct ctgctcttct tctacctttc aagtttttaa agtattaaga 120  
tggcagagac attcctatctt acctcagagt cagtgaacga gggacaccct gacaagctct 180  
gcgaccaaatt ctccgatgct gtcctcgacg cttgccttga acaggaccca gacagcaagg 240  
ttgcctgcga aacatgcacc aagaccaact tggatcatgg ctccggagag atcaccacca 300  
nggc 304

<210> 2055  
<211> 481  
<212> nucleic acid  
<213> Glycine max  
<400> 2055

aaactccacc gccaggtac cggatcaaga attcccggtt cgaccacgc gtcngggcag 60  
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acagaaggggt acggctgcga agaagacgac agaaggggtac ggctgcgaga agacgacaga 180  
aggggtacggc tgcgagaaga cgacagaang gtacggctgc gagaagacga cagaaggggg 240  
acacttatgg aactgggaag attcctgaca aggagattct tcaaattgtg aaggagaatt 300  
tcgacttcag acctggaatg atcaccatta acttgacct taagaggggt ggccataggt 360  
tctcaagac agctgcttat ggacactttg gaagggatga ccctgacttc acctgggaag 420  
ttgtgaagcc actcaantct gaaaaacctc caacctaga atggttgtna atttaancnc 480  
c 481

<210> 2056

<211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2056  
  
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 gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ctttgccaga 120  
 agttaaaatg gcccaagaaa ctttcttatt cacatctgaa tcagtgaacg aggggcaccc 180  
 tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttcg agcaggaccc 240  
 tgacagcaag gttgcctnt gaaacctgca ccaagaccaa catggtgatg gttttcggag 300  
 agatcacaac caa 313

<210> 2057  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2057  
  
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 tcctattcac atctgaatca gtgaacgagg ggcacctga caagctctgt gaccagatct 180  
 ccgatgctgt gctcgatgca tgcttgagc aggacctga cagcaagggt gcctgtgaaa 240  
 cctgacacca agaccaacat ggtgatgggt ttccggagaga tcacaaccaa ggccaacgtg 300  
 gatatg 306

<210> 2058  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2058  
  
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 gggttactgtc tgttcaagct accatctctc tctctctttt ttagtgcctc cttgccagaa 120  
 gttaaaatgg cccaaganac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
 gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttga gcaggaccct 240

gacagcaagg ttgcctgtga aacctggcac caagaccaac atggtgatgg ttttcggaga 300  
 gatcacaacc aaggccaagt ggata 325

<210> 2059  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2059

tcgcatgcac gcgtacgtna gctcggaatt cggctcgagc tttctgctct tttctacct 60  
 ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag antcggtgaa 120  
 cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt 180  
 cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggttca 240  
 tgggtcttcgg agaaatcacg accaaggcca acgttgacta cgagaa 286

<210> 2060  
 <211> 280  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2060

gtcgcangca cgcgtacgta aagctcggaa ttcggctcga gnaaagatgg cagagacatt 60  
 cctatattacc tcagagtcgg tgaacgaggg acaccctgac caagctctgc gaccaaattct 120  
 ccgatgctgt cctcgacgct tgctcgcagc aggncccaga tagcaaagtt ncntgcgana 180  
 catgcaccan aaccnncttg gtcattggtct tcggagnnat cagcaccang gcnancgttg 240  
 actanganan gatantgngt gacacctnca ggnacatcgg 280

<210> 2061  
 <211> 324  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2061

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 actgaactaa tcattaattt gcactcgcgtg tttcagcttc atcacccttc ttttgcattca 120  
 tttatatctc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180

agggtcaccc cgacaagctg tgnaccaga tctctgatgc agtacttgat gcgtgccttg 240  
ancaggaccc tgacagcaag gttgcctgtg agacatgcac cnagaccaac aggtcatggt 300  
cttcggagag atcacaacca aggc 324

<210> 2062  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2062

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gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca gaagttaaaa 120  
tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180  
tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240  
aggttgcttg tgaaacctgc accaagacca acatggtgat ggtttcggag agatcacaac 300

<210> 2063  
<211> 227  
<212> nucleic acid  
<213> Glycine max  
<400> 2063

ntcgcanaca cgcgtacgtn agncgggaat tcggctcgag gtggcaaattg gccttnccag 60  
aaggtgcatt gtccaagttt cctatgccat tgggtgccct gagcccttgt cagtgtttgt 120  
ggacacttat ggaactggga agattcctga caaggagatt cttcaaattg tgaaggagaa 180  
tttcgacttc agacctggaa tgatcaccat taacttggn c ttaaann 227

<210> 2064  
<211> 313  
<212> nucleic acid  
<213> Glycine max  
<400> 2064

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acntctcttc gttcacgcta cccctttctg gctcttcttc tacctttcaa gttttaaaag 120  
tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacaccctga 180



caagctctgc gaccaaactc cccgatgctgt cctcgacgct tgcctcgagc aggacccaga 240  
cagcaaagtt gcttgcgaaa catgcaccaa aaccaacttg gtcattgtct tcggagaaat 300  
cacgaccaag gcc 313

<210> 2065  
<211> 311  
<212> nucleic acid  
<213> Glycine max

<400> 2065

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ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180  
caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag 240  
gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaaga ccaacatggg gatggttttc 300  
ggagagatca n 311

<210> 2066  
<211> 317  
<212> nucleic acid  
<213> Glycine max

<400> 2066

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agctctgcga ccaaactctc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240  
gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt catggtcttc ggagaaatca 300  
cgaccaaggc caacgtt 317

<210> 2067  
<211> 306  
<212> nucleic acid  
<213> Glycine max

<400> 2067

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 aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180  
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240  
 ctgacagcaa ggttgccctgt gaaacctgca ccaagaccaa catggtgatg gttttcggag 300  
 agatca 306

<210> 2068  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max

<400> 2068

ancagtcgna tgcacgcgta cgtaagctcg gaattcgget cgagccccac tcaaccacca 60  
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 tataaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 180  
 caagctctgc gaccaaattc ccgatgctgt cctcgacgct tgccctcgagc aggaccacaga 240  
 cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatggtct tcggagaaat 300  
 cacgaccaag gccaaagtga 320

<210> 2069  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max

<400> 2069

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 gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg 180  
 acaagctctg cgaccaaatt tccgatgctg tcctcgacgc ttgcctcgag caggaccacag 240  
 acagcaaagt tgctgcgaaa acatgcacca aaaccaactt ggtcatgggtc ttggagaaaa 300  
 tcacgaccaa ggccaagt 318

<210> 2070

<211> 302  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2070  
  
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 aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180  
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240  
 ctgacagcaa ggttgccctgt gaaacctgca ccaagaccaa catggtgatg gttttcggag 300  
 ag 302

<210> 2071  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2071  
  
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 ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg taaacgaggg 180  
 tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctcgatgcgt gccttgaaca 240  
 ggaccctgac agcaaggttg cctgtgagac atgcaccaag accaacaatgg tcatggtc 298

<210> 2072  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2072  
  
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 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240  
 ccctgacagc aaggttgccct gtgaaacctg caccaagacc aacatggtga tggttttcgg 300

agagatcacn 310

<210> 2073  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2073

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 aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc tctgtgacca 180  
 gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca aggttgccctg 240  
 tgaaacctgc accaagacca acatggtgat ggttttcggg gagatcaca 289

gagatcacn

<210> 2074  
 <211> 309  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2074

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 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240  
 ccctgacagc aaggttgccct gtgaaacctg caccaagacc aacatggtga tggttttcgg 300  
 agagatcac 309

<210> 2075  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2075

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 gaagttaaaa tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac 180

cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac 240  
cctgacagca aggttgcttg tgaaacctgc accaagacca acatgggtgat ggtttttcgga 300  
gagatcac 308

<210> 2076  
<211> 310  
<212> nucleic acid  
<213> Glycine max  
<400> 2076

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gttaaaatgg cccaaganac tttcctattc acntctgant cngtgaacga ggggcaccct 180  
gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
gacagcaagg ttgctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300  
atcacaacca 310

<210> 2077  
<211> 310  
<212> nucleic acid  
<213> Glycine max  
<400> 2077

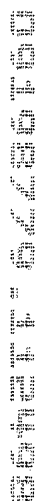
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aactcgttc atatatctct ctgctcttct cttactcttc tacctctcaa gtttttgaag 120  
tataaagatg gcagagacat tcctattcac ctcgagtgca gtgaacgagg gacaccctga 180  
taagctctgc gaccaaattc ccgatgctgt cctcgacgct tgctcgaac aggaccaga 240  
cagcaagggt gctgcgaaa catgcacca gaccaacttg gtcatggtct tcggagagat 300  
caccaccaag 310

<210> 2078  
<211> 325  
<212> nucleic acid  
<213> Glycine max  
<400> 2078

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 gccanaagtt aaaatggccc aagaaacttt cctatncaca tctgaatcag tnaacgangg 180  
 gcaccctgac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gctgggagca 240  
 ggaccctgnc agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatggtttt 300  
 cggagagatc acaaccaagg nnagc 325

<210> 2079  
 <211> 249  
 <212> nucleic acid  
 <213> Glycine max

<400> 2079



ctcgagccgc tcgagccgat tcggctcgag ctcccagact catncccttg agccatgtca 60  
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 tgagacctga tggcaagacc caagtcactg ttgagtacta caatgacaag ggtgccatgg 180  
 ttccaatccg cgtccacact gtgctcatct ccacacagca tgatganctg tcacaaatga 240  
 tgagattgc 249

<210> 2080  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max

<400> 2080

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 caaagcgggt tactgtctgt tcaagctacc atctctctct ctctttotta gtgcctcctt 120  
 gccagangtt naaatgggcn caagaaactt tcctattcac atctgnatca gtgaacgagg 180  
 ggcaccctga caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttgagc 240  
 aggaccctga cagcaagggt gcctgtgaaa cctggcacca agaccaacat ggtgatgggt 300  
 ttoggagaga tcacaaccaa ggcca 325

<210> 2081  
 <211> 316  
 <212> nucleic acid

<213> Glycine max

<400> 2081

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gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ctttgccaga 120  
agttaaaatg gcccaagaaa ctttcttatt cacatctgaa tcagtgaacg aggggcaccc 180  
tgacagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300  
atcacaacca aggcca 316

<210> 2082

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2082

ncgtcgcatg cagcggtacg taagctcggg atttcggctc gaggggttac tgtctgttca 60  
agctaccatc tctctctctc tttcttagtg ctccttgcc agaagttaa atggccaag 120  
aaacttttct attcacatct gaatcagtga acgaggggca ccctgacaag ctctgtgacc 180  
agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aaggttgcct 240  
gtgaaacctg cnaccaagac caacatggtg atggttttcg gagagatcac aaccaangcc 300  
aac 303

<210> 2083

<211> 333

<212> nucleic acid

<213> Glycine max

<400> 2083

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag aagccccact caaccaccac 60  
acctctctc gttcacgcta cccctttctg ctcttcttct acctttcaag ttttaaaagt 120  
ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgagg acacctgac 180  
aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca ggacctgac 240  
agcaaagttg cctgcgaaac atgcacaaa accaattggt catggtcttc ggagaaatca 300

cgaccaaggc caagttgact acgagaagat atg

333

<210> 2084  
<211> 287  
<212> nucleic acid  
<213> Glycine max

<400> 2084

gcacgcgtac gtaagctcgg aattcggctc gaggtgcctt ctctgggaag gatcctacca 60

aggttgatag gagtgggtgcc tacattgtga ggcaagctgc aaagagcatt gttgcaaatg 120

gacttgctag gagggcaatt gtgcaagttt cctatgccat tgggtgtgcct gagccctgtc 180

tgtgtttgtt gacaattatg gcaactggga gatcccgaca aggaaatcct cagcatgtga 240

aggagagttt tgaactcagcc ggcagatctc catcaacctg atctcaa 287

<210> 2085  
<211> 281  
<212> nucleic acid  
<213> Glycine max

<400> 2085

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accatctctc tctctctttc ttagtgcctc cttgccagaa gttaaaatgg cccaagaaac 120

tttcttatte acatctgaat cagtgaacga ggggcacctt gacaagctct gtgaccagat 180

ctcogatgct gtgctcgatg catgcttgga gcaggacctt gacagcaagg ttgcctgtga 240

aacctgcacc aagaccaaca tggatgatgg tttcggagag a 281

<210> 2086  
<211> 294  
<212> nucleic acid  
<213> Glycine max

<400> 2086

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gttactgtct gttcaagcta ccatctctct ctctctttct tagtgcctcc ttgccagaag 120

ttaaaatggc ccaagaaact ttcctattca catctgaatc agtgaacgag gggcaccttg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggacctg 240



acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggtt ttcg 294

<210> 2087  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max

<400> 2087

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ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcacctg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggacctg 240

acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggtt ttcg 294

<210> 2088  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max

<400> 2088

nngtcgancg cagcgtacg taagctcgga attcggctcg agacagcaca aagcgggtta 60

ctgtctgttc aagctaccat ctctctctct cttcttagt gcctccttgc cagaagttaa 120

aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc accctgacaa 180

gtctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240

caaggttgcc tgtgaaacct gcaccaagac caacatgggt atggttttcg 290

<210> 2089  
 <211> 322  
 <212> nucleic acid  
 <213> Glycine max

<400> 2089

agtcgcangc angcgtacgt nagctcgga ttcggctcga ggcagactta acaacagcac 60

aaagcgggtt actgtctgtt caagctacca tctctcctc tctttcttag tgctccttg 120

ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gnacgagggg 180

cacctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg ctcggagcag 240

gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaagn ccaacntggt gatggttttc 300  
ggagannnca anccaagggc an 322

<210> 2090  
<211> 318  
<212> nucleic acid  
<213> Glycine max  
<400> 2090

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accaccacac ctctctctgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120  
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240  
accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300  
gagaaatcac gaccaagg 318

<210> 2091  
<211> 301  
<212> nucleic acid  
<213> Glycine max  
<400> 2091

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gttactgtct gttcaagcta ccattctctc ctctctttct tagtgcttcc ttgccagaag 120  
ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcaccctg 180  
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccctg 240  
acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgggt ttcgagagaga 300  
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<210> 2092  
<211> 289  
<212> nucleic acid  
<213> Glycine max  
<400> 2092

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acacnactct ctctgctctt cttctacctt tcaagttttt aaagtattaa gatggcagag 120  
acattcctat ttacctcaga gtcagtgaac gagggacacc ctgacaagct ctgcgaccaa 180  
atctccgatg ctgtcctcga cgcttgccctt gaacaggacc cagacagcaa ggttgccctgc 240  
gaaacatgca ccaagaccaa cttgggtcatg gtcttcggag agatcacca 289

<210> 2093  
<211> 309  
<212> nucleic acid  
<213> Glycine max  
<400> 2093

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aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa 180  
gtcttgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 240  
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gaccaaggc 309

<210> 2094  
<211> 336  
<212> nucleic acid  
<213> Glycine max  
<400> 2094

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accaccacac ctncgcctcg ttcacgctac ccctttctgc tcttcttcta cctttcaagt 120  
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg 180  
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240  
gaccagaca gcaaagttgc ctgcgaaaca tgcaccanaa ccaacttggt catggtcttc 300  
ggagaaatca cgaccaaggc caagttgact acgaga 336

<210> 2095  
<211> 202  
<212> nucleic acid  
<213> Glycine max

<400> 2095

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atgcttggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
ggtgatgggtt ttgggagaga tc 202

<210> 2096

<211> 315

<212> nucleic acid

<213> Glycine max

<400> 2096

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctcgagccgg aacttaacaa 60  
cagcacaaaag cgggttactg tctgttcaag ctaccatctc tctctctctt tottagtgcc 120  
tccttggcag aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac 180  
gagggggcacc ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgctt 240  
gagcaggacc ctgacagcaa ggttgccctgt gaaacctgca ccaagaccaa catggtgatg 300  
gttttcggag agatc 315

<210> 2097

<211> 322

<212> nucleic acid

<213> Glycine max

<400> 2097

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tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg 180  
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240  
gaccagaca gcaaagttgc ctgcgaaaca tgcaccanna ccaacttggc catggtcttc 300  
ggagaaatca cgaccaaggc ca 322

<210> 2098

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 2098

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accacaccac tctntctggc tcttcttcta ctttcaagt tnttaaagta ttaagatggc 120  
ngagacagcc ctatttacn cagagtcagt gaacgangga caccctgaca agctctgcga 180  
ccaaatctcc gatgctgtcc tcgacgcttg ccttgaacag gaccagaca gcaaggttgc 240  
ctgcgaaaca tgcaccaaga ccaacttggc catggtctnc ggagagatca ccaccaaggc 300  
caacgtt 307

<210> 2099

<211> 323

<212> nucleic acid

<213> Glycine max

<400> 2099

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ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
acaccctgac aagctctgag accaaatctc cgatgctgtc ctcgacgctt gcctcgagcn 240  
cgaccagac agcaaagttg cctgcgaaac atgcaccaan accaacttgg tcatggtctt 300  
cggagaaatc acgaccaagg cca 323

<210> 2100

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2100

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ccagcgagtg ttctttcttc gtttcaacac cttaatttgc acacgctgct tcttcagctt 120  
gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag ggtcaccctg 180  
acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa caggaccctg 240  
acagcaaggt tgctgtgag acatgcacca agaccaacat ggtcatggt 289



gaaacatgca ccaagaccaa cttgggtcatg gtcttcggag agatcaccac caaggccaag 300  
 ttgactagag a 311

<210> 2104  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2104

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 agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
 cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac 240  
 cctgacagca aggttgcttg tgaaacctgc accaagacca acatgggtgat ggtttttcgga 300  
 gagatcacia cca 313

<210> 2105  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2105

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 tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180  
 aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240  
 agcaagggtg cctgtgaaac ctggcaccaa gaccaacatg gtgatggttt tcggagagat 300  
 cacaac 306

<210> 2106  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2106

agttcanaca gcgtacgana gctcggaant cggctcgagg gccaggcnag cccnatcaac 60

cancacacnt ctctacnct cacgctacnc cttgctgcnc ttncgcgac ntngcaagt 120  
nctnaaaagt ataaagatgg cagagacatn cctantnacc ncagagtcgg tgaacgaggg 180  
anaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240  
ggacccagac agcaaagttg cctgcgaaac atgcaccaa accaacttgg tcatggtctt 300  
cggagaaatc acgaccaagg ccaac 325

<210> 2107  
<211> 294  
<212> nucleic acid  
<213> Glycine max  
<400> 2107

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ctgtctgttc aagctaccat ctctctctct tttcttagt gcctccttgc cagaagttaa 120  
aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc accctgacaa 180  
gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240  
caaggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttcg gaga 294

<210> 2108  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
<400> 2108

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agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120  
agaagttaaa atggcccaaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
cctgacaaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240  
cctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300  
agan 304

<210> 2109  
<211> 303  
<212> nucleic acid  
<213> Glycine max



<400> 2109

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agcgggttac tgtctgttca agctacnate tctctctctc tttcttagtg cctccttgcc 120

agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180

ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240

ccctgacagc aaggttgctt gtgaaacctg caccaagacc aacatggtga tggttttcgg 300

aga 303

<210> 2110

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2110

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gtttctgacg ctctgtcttc agcgagtgtt ctttcttctg ttcaaacacct taatttgac 120

acgtgtcttc ttacagcttga gaantggcac aagaaacctt tctattcaca tctgaatctg 180

taaacgaggg tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctgatgcgt 240

gccttgaaca ggaccctgac agcaagggtt cctgtgagac atgcaccaag accaacaagg 300

tca 303

<210> 2111

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2111

acgtcgcattg cacgcgtacg taagctcggaa attcggctcg agatttgga gttaggttct 60

gcacgctctg cttccagcga gtgttctttc ttcgtttcaa caccttaatt tgcatacgt 120

gcttcttcng cttgagaaat ggcaagaagaa acctttctat tcacatctga atctgtaaac 180

gagggtcacc ccgacaagct gtgcgaccag atctctgatg cagtgcctga tgcgtgcctt 240

gaacaggacc ctgacagcaa ggttgctctg gagacatgca ccaagaccaa catggtca 298

<210> 2112

<211> 286  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2112  
  
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 ccaactctctc tgctcttctt ctacctttca agtttttaaa gtattaagat ggcagagaca 120  
 ttctatttta cctcagagtc agtgaacgag ggacaccctg acaagctctg cgaccaaatac 180  
 tccgatgctg tnnctgacgc ttgccttgaa caggaccag acagcaaggt tgcttgcgaa 240  
 acatgcacca agaccaactt ggtcatggtc ttccggagaga tcacca 286

<210> 2113  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2113  
  
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 aaattgtgaa ggagaatttc gacttcagac ctggaatgat caccattaac ttggacctta 120  
 agaggggtgg tcataggttc ctcaagacag ctgcttatgg acactttgga agggatgatg 180  
 cagacttcac ctgggaagtt gtgaagccac tcaagtcaga gaagcctcaa gcttaagagt 240  
 gttgttaagt taatcactcc cttcagtggg tgtcttgctg ggtgtggatg aataatttgc 300  
 gtgtttcatg actact 316

<210> 2114  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2114  
  
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 gaagtacott gatgacatgc tctttaaggt ccgcagcaat ttgatcattg ctcacagtct 120  
 ncatcatggt ggggtggacct taaagagcan nttntcaagc ctgtcattcc tgagaagtac 180  
 cttgatgaga agaccatctt ccaccttaac ctttctggcc gttttgtcat tgggtggcct 240  
 catggtgang ctgcnctcac tggaagaaa atcatcattg atacctatgg tggctggggg 300

gctcatgg

308

<210> 2115  
<211> 284  
<212> nucleic acid  
<213> Glycine max

<400> 2115

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caenactctc tctgctcttc ttctaccttt caagttttta aagtattaag atggcagaga 120  
cattcctatt tacctcagag tcagtgaacg agggacaccc tgacaagctc tgcgaccaa 180  
tctccgatgc tgtctcgac gcttgcttg aacaggaccc agacagcaag gttgctgcg 240  
aaacatgcac caagaccaac ttggtcatgg tcttcggaga gatc 284

<210> 2116  
<211> 283  
<212> nucleic acid  
<213> Glycine max

<400> 2116

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accactctct ctgctcttct tctacctttc aagtttttaa agtattaaga tggcagagac 120  
attcctatct acctcagagt cagtgaacga gggacaccct gacaagctct gcgaccaa 180  
ctccgatgct gtctctgacg cttgcttga acaggacca gacagcaagg ttgctgcca 240  
aacatgcacc aagaccaact ttggtcatgg cttcgagag atc 283

<210> 2117  
<211> 298  
<212> nucleic acid  
<213> Glycine max

<400> 2117

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gcggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcttgcca 120  
gaancgcaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240

ccctgacagc aaggttgcc t gtaaacctg caccaagacc aacatgggtga tggttttc 298

<210> 2118  
<211> 288  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2118

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ttccagcgag tgttctttct tcgtttcaac accttaattt gcacacgctg cttcttcagc 120  
ttgagaaatg gcacaagaaa ctttctatt cacatctgaa tctgtaaagc agggtcaccc 180  
cgacaagctg tgcgaccaga tctctgatgc agtgcctgat gcgtgccttg aacaggaccc 240  
tgacagcaag gttgcctgtg agacatgcac caagaccaac atgggtcat 288

<210> 2119  
<211> 329  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2119

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acacctctcn tcgttcangc tannnaaatn ctgctgttct tctacctgac aagttttgaa 120  
agtatanaga tggcaganac attcctattt acctcanagt cggatgaacga gggacaccct 180  
gacaagctct gcgaccaaatt ctccgntgct gtccctcgacg cttgcntcga gcagnaccca 240  
gacagcaaag ttgccngcga nacatggacc aaaaccaact tggatcatgg ntccggagaa 300  
atcacgacca aggccaaact tgactacnn 329

<210> 2120  
<211> 277  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2120

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cttctttctac ctttcaagtt ttaaaagtat aaagatggca gagacattcc tatttnoctc 120  
agagtcggtg aacgaggagc accctgacaa gctctgcgac caaatctccg atgctgtcct 180

cgacgcttgc ctcgagcagg acccagacag caaagttgcc tgcgaaacat gcacccaaaac 240  
 caacttggtc atggtcttcg gagaaatcac gaccaag 277

<210> 2121  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2121

cgcangcacg cgtacgtaag ctcggaattc ggctcgagca agccccactc aaccaccaca 60  
 cgcgtenctc tngcgcttct tctaccttct aagtttttaa agtattaaga tggcaganac 120  
 attcctatctt acctcagagt cagtgaacga gggacaccct gacaagctct gcgaccaaact 180  
 ctccgatgct gtctcgcagc cttgccttga acaggaccca gacagcaagg ttgcctgcga 240  
 aacatgcacc aagaccaact tggatcatggt cttcggagag atcacc 286

<210> 2122  
 <211> 339  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2122

annctgaanc gtangnaagc ntacgnattc ngctcgagng gcaggcaagc cccactcaan 60  
 caccacacct gctcctgcgt ncangctnac ccgtnnnngan gnatgacta cctntcaagt 120  
 tntaaaagta tngnanatgg cngagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
 aacccctgac aagctctgcg accaaatctc cgntgctgtc ctcgacgctt gcctcgagca 240  
 ggaccagac agcaaagttg cctgcgaaac atgcaccacc accaagttgg tcatggtctt 300  
 cggagaaatc acgaccaagg cnaacgttac tacgagann 339

<210> 2123  
 <211> 480  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2123

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 gtacggctgc gagaagacga cagaaggggg cagctcttga tttnaggnca ngcaancccc 120

actcaanac cacacctctc ctcggtcacg ctatcccttt ctgctcttct tctacctttc 180  
angttttaan agtacncaca tggcaagaca cattcctatt tancnagac tcggtgaann 240  
acggacaccc tgacaagctc tgcgaccaa tctccgatnc tgtcctcgac gcttgccctcg 300  
ancaggactc agacancana nttgcctgcn aaacatgcac caaaaccaac ttggtcatgg 360  
tcttcngaga antcacgacc aaggccaach ttgactacga aaaganngtg cgttacacct 420  
gcgggaaca tcggcttctt tcnaaatgat gttgggactg gatgccgacc actgcatngg 480

<210> 2124  
<211> 307  
<212> nucleic acid  
<213> Glycine max

<400> 2124

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gggggttaact gtctgttcaa gctaccatct ctctctctac tttcttagtg cctccttgcc 120  
agangttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
ccctgacaag ctctgtgacc agatctccga tgcgtgtgctc gatgcatgct tggagcagga 240  
nnctgacag caagggttgcc tgtgaaacct gcaccaagac caacatggtg atggttttcg 300  
gagagat 307

<210> 2125  
<211> 307  
<212> nucleic acid  
<213> Glycine max

<400> 2125

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gttactgtct gtccaagcta ccatctctct ctcnctttc ctnagtgcct ccttgccaga 120  
agttaaaatg gcccaagaaa ctttcttatt cacatctgaa tcagtgaacg aggggcaccc 180  
tgaaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
gacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgg tttcggagag 300  
atcacia 307

<210> 2126

<211> 309  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2126  
  
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 ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa 120  
 aatggcccaa gaaactttcc tattcacatc tgaatcagt aacgaggggc accctgacaa 180  
 gctctgtnac cagatctccg atgctgtgct cgatgcatgc ttggagcagg accctgacag 240  
 caaggttgcc tgtgnaaacc tggcaccaag accaacaatgg tgatggtttt cggagagatc 300  
 acaaccaag 309

<210> 2127  
 <211> 302  
 <212> nucleic acid  
 <213> Glycine max  
  
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 atctctctgc tcttctcttc tcttctacct ctcaagtttt tgaagtataa agatggnaga 120  
 gacattccta ttcacctcgg agtcagtga cgagggacac cctgataagc tctgcgacca 180  
 aatctccgat gctgtcctcg acgcttgctt cgaacaggac ccaganagca aggttgccctg 240  
 cgaaacatgc accaagacca attggatcatg gtcttcggag agatcaccac caaggccaac 300  
 gt 302

<210> 2128  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
  
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 cccaagaaac tttcctattc acgtctgaat cagtgaacga ggggcaccct gacaagctct 180  
 gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240





accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

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<210> 2132

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2132

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aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120

cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180

accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

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<210> 2133

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2133

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gaaggagaat ttcgacttca gacctggaat gatcaccatt aacttggacc ttaagagggg 120

tggccatagg ttcttcaaga cagctgctta tggacacttt ggaagggatg accctgactt 180

cacctgggaa gttgtgaagc cactcaagtc tgagaagcct caagcttaag attgttgtga 240

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atgactacta ctacttcac 320

<210> 2134

<211> 313

<212> nucleic acid

<213> Glycine max

<400> 2134

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caccacacct ctctctgttc acgctacccc tttctgctct tcttctacct ttcaagtttt 120

aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa cgagggacac 180  
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240  
 ccagacagca aagttgcctg cgaaatntgc accaaaacca acttggtcat ggtcttcgga 300  
 gaaatcacga cca 313

<210> 2135  
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 <212> nucleic acid  
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 ngtttttaaaa gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag 180  
 ggacaccctg acaagctctg cgaccaaate tccgatgctg tcctcgacgc ttgcctcgag 240  
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 ttccggagaaa tcacga 316

<210> 2136  
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 <213> Glycine max  
 <400> 2136

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 aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180  
 ctgacaagct ctgogaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240  
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 aaatcacga 309

<210> 2137  
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<400> 2137

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aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccngac aagctctgtg 180  
accagatctc cgatgctgtg ctcgatgcat gcttgagca ggaccctgac agcaagggtg 240  
cctgtgaaac ctgcaccaag accaacaatgg tgatggtttt 280

<210> 2138

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2138

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aggttctgca cgctctgctt ccagcgagtg ttctttcttc gtttcaacac ctttaatttg 120  
acaagctgct tcttcagctt gagaaatggc acaagaaacc tttctattca catctgaatc 180  
tgtaaacgag ggtcaccctg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc 240  
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ggt 303

<210> 2139

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2139

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cgctctgctt ccagcgagtg ttctttcttc gtttcaacac ctttaatttg acacgctgct 120  
tcttcagctt gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag 180  
ggtcaccctg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa 240  
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<210> 2140

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 2140

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ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa 180  
cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct 240  
cgagcaggac ccagacagca aagtngcctg cgaaanatgc accagaacca acttggtcat 300  
ggtcttcgga gaaatcacga ccaag 325

<210> 2141

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2141

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attgtgaagg agaattncga cttcanacct ggaatgatca ccattaactt ggaccttaag 120  
aggggtggcc ataggttcct caagacagct gcttatggac actttggaag ggatgaccct 180  
gaattcacct gggaagttgt gaagccactc aagtctgaga agcctcaagc ttaagattgt 240  
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<210> 2142

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2142

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gttctgcacg ctctgcttcc agcgagtgtt ctttcttcgt ttcaacacct taatttgcac 120  
acgctgcttc ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180  
taaacgaggg tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctcgatgcgt 240  
gccttgaaca ggaccctgac agcaaggttg cctgtgagac atgcaccaag accaacaatgg 300  
t 301

<210> 2143  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2143  
  
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 agctaccatc tctctctctc tttcttagtg ctccttgcc agaagttaa atggcccaag 120  
 aaactttcct attcacatct gaatnagtga acgaggggca ccctgacaag ctctgtgacc 180  
 agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc aaggttgcct 240  
 gtgaaacctg caccaagacc aacatggtga tggttttcgg aga 283

<210> 2144  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2144  
  
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 tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180  
 aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240  
 agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatgggttt cgg 293

<210> 2145  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2145  
  
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 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180  
 aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240  
 agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatgggttt cgga 294

<210> 2146  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2146  
  
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 gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga 120  
 agttaaaatg gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc 180  
 tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc 240  
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atgggtgatgg t 291

<210> 2147  
 <211> 340  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2147  
  
 acgogtacgt aagctcgga ttcggctcga gggccaggca agccccactc aaccaccaca 60  
 cntctectgc gttcangcta cccctttctn gctcttcttc tacctntcaa gtnttaaaag 120  
 tataaagatg gcagagacat tctattttac ctcagagtcg gtgaacgagg gacaccctgn 180  
 caagctctgc gaccaaatct ccgatgctgt cctcgacgct tgccctcgagc aggaccacaga 240  
 cagcaaagtt gcttggcgaa acatgcacca ntnnnacttg gtcatgggtct tcggagaaat 300  
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<210> 2148  
 <211> 319  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2148  
  
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 cactcaacca ccacacctct cctcgttcac gctaccctt tctgctcttc ttctaccttt 120  
 caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180  
 agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgcctcg 240

agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaaccaac ttggtcatgg 300  
tcttcggaga aatcacgac 319

<210> 2149  
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tagtgctcc ttgccagaag ttaaaatggc ccaaganact ttctattca natctgaatc 60  
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atgcttgag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
ggatgatggt ttccgaga 198

<210> 2150  
<211> 293  
<212> nucleic acid  
<213> Glycine max  
<400> 2150

ngtencangc acgcgtacgt nagctcgga ttccgctcga ggcacaaagn gggttactgt 60  
ctgttcaagc taccatctct ctctctgctt tgcttagtgc ctccctgcca gaagttaaaa 120  
tgccccaaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180  
tctgtgacca gatctccgat gctgtgctcn ngccatgctt ggagcaggac cctgacagca 240  
aggttgcntg tgaaacctgc accaagacca acatggatgat ggttttcgga gag 293

<210> 2151  
<211> 295  
<212> nucleic acid  
<213> Glycine max  
<400> 2151

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cgtctcaa at gatgtgggac tggatgccga caactgcaag gtcctcgtca acattgagca 180  
gcagagccnt gatattgcct cagggtgtac acggnccacc ttacnnnnaa acctgaagaa 240

nttgggtgctg gtgaccaggg tccacatggt tggctatgcc atgatgaaac cccnc 295

<210> 2152  
<211> 219  
<212> nucleic acid  
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<400> 2152

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agtgaacgag gggcaccctg acaagtctgt gaccagatct ccgatgctgt gctcgatgca 120  
tgcttgagc aggaccctga cagcaaggtt gctgtgaaa cctggcacca agaccaacat 180  
ggtgatggtt ttgggagaga tcacaancaa ggccaacgt 219

<210> 2153  
<211> 218  
<212> nucleic acid  
<213> Glycine max

<400> 2153

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agtgaacgag gggcaccatc gacaagctct gtgaccagat ctccgatgct gtgctcgatg 120  
catgcttggc gcaggaccct gacagcaagg ttgctgtga aacctgcacc aagnaccaac 180  
atggtgatgg ttttcggaga gatcacaacc aaggccaa 218

<210> 2154  
<211> 291  
<212> nucleic acid  
<213> Glycine max

<400> 2154

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aagcnaccat ctncctctct ctttcttagt gctccttgc cagaagttaa aatggcccaa 120  
gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa gctctgtgac 180  
cagatctccg atggctgtnc tcgatgcatg cttggagcag gacctgaca gcnaggttgc 240  
ctgtgaaacc tgcaccaaga ccaacatggt gatggttttc ggagagatca n 291

<210> 2155



<211> 309  
 <212> nucleic acid  
 <213> Glycine max  
  
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 ccagaagtta aaatggggccc aagaaacttt cctattcaca tntgaatcag tgaacgaggg 180  
 gcaccctgac aagctctgtg accagatctc cgatgctgtg cttcgatgca tgcttggagc 240  
 aggaccctga cagcaagggtt gcctgtgaaa cctgcaccaa gaccaacatg gtgatgggtt 300  
 tcggagaga 309

<210> 2156  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2156  
  
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 nataaagatg gcagagacat tcctatttan ctcagagtcg gtgaacgagg gacaccctga 180  
 caagctctgc gaccaaactc ccgatgctgt cctcgacgct tgccctcgagc aggaccaga 240  
 cagcaaagtt gcctgcgana catgccacca aaaccaactt ggtcatggtc ttcgagaaaa 300  
 tcacgaccaa ggc 313

<210> 2157  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max  
  
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 ttcacctcgg agtcagtga cagaggacac cctgataagc tctgcgacca aatctccgat 180  
 gctgtcctcg acgcttgctt cgaacaggac ccagacagca aggttgcttg cgaaacatgc 240

accaagacca acttggttca tgggtcttcgg agagatcacc accaaggcca acgt 294

<210> 2158  
 <211> 285  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2158

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 ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat ggcccaagaa 120  
 acttttctat tcacatctga atcagtgaac gaggggcacc ctgacaagct ctgtgaccag 180  
 atctccgatg ctgtgctcga tgcattgctt gagcaggacc ctgacagcaa ggttgccctgt 240  
 gaaacctggc accaagacca acatggtgat gggtttcggg gagat 285

CCAGGTTCA TGGGTCTTCGG AGAGATCACC ACCAAGGCCA ACGT

<210> 2159  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2159

gcacgngtac gtnagctcgg aattcggctc gaggccattt gggagttagg ttctgcacgc 60  
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 tcagcttgag aaatggcaca agaaaccttt ctattcacat ctgaatctgt aaacgagggg 180  
 caccctgaca agctgtgcga ccagatctct gatgcagtgc tcgatgcgtg ccttgaacag 240  
 gacctgaca gcaaggttgn ctgtgagact gcaccaagac caacatgggc atggtctttg 300

<210> 2160  
 <211> 258  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2160

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 gccaacgat gaaacccccg agtaactgcc ctcagccat nctccttgca accaaacttg 120  
 gngctcgnt cacagaggtt aggaagaatg gcacctgtgc ttggttgagg ccagatggta 180  
 agacacaagt aaccgtcgag tactacaatg acaatgggtc catggttcca gttcgtgtcc 240

acactgtcct aatttcca

258

<210> 2161  
<211> 335  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2161

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agccccactc aacnaccaca cctctcctcg ttcacgctac ccctttctgc tcttcttcta 120  
ccttcaagtt ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcgggtg 180  
aacgagggac accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc 240  
ctcgagcagg acccagacag caaagttgcc tgcgaaacat gcacaaaaac caattgggtca 300  
tggctcttcg agaaatcacg accaaggcca acgtt 335

<210> 2162  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2162

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accactctct ctgctcttct tctaccttct aagtnggtaa agtattaaga tggcagagac 120  
attcctatctt acctcagagt cagtgaacga gggacaccct gacaagctct gcgaccaaatt 180  
ctccgatgct gtctctgacg cttgccttga acaggaccca gacagcaagg ttgcctgcga 240  
aacatgcacc aagaccaact tggctcatggt cttggagaga tcaccac 287

<210> 2163  
<211> 319  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2163

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acagcacaaa gcgggttact gtctgttcaa gntanccatc tntgctctct ctttcttagt 120  
gcctccttgc nagaagntan aatggcccaa gnaaactttc ctattcacat ctgaatcagt 180

gancgagggg caccctgaca agctctgtga ncagatctcc gatgctgtgc tcgatgcatg 240  
 cttggagcag gaccctnaca gcaaggttgc ctgtgaaacc tgcaccaaga ccaanatggt 300  
 gatngttttc ggagagatc 319

<210> 2164  
 <211> 327  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2164

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 tgtncaaagt accattctct ctctctcttt cttagtgcct ccttgccata agttaaaatg 120  
 gcccnagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc tgacaagctc 180  
 tgtgaccaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc tnacagcaag 240  
 gttgcctgtg aaacctgcac caagaccaac atggtgatgg tttcggagag atcacgacca 300  
 aggncaantg ggtntgagaa gatngtg 327

<210> 2165  
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 <212> nucleic acid  
 <213> Glycine max  
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 accaccacac ctctctctgt tcacgctacc ctttctgtct cttctctac ctttcaagtt 120  
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctgagcagg 240  
 acccagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300  
 gagaaatca 309

<210> 2166  
 <211> 260  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2166

aagccccact caaccaccac acctctctc gttcacgcta cccctttctg ctctttctct 60  
 acctttcaag ttttaaaagt ataaagatgg cagagacatt cctattttacc tcagagtcgg 120  
 tgaacgaggg acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt 180  
 gcctcgagca ggaccagac agcaaagttg cctgcgaaac atgcaccaa accaacttgg 240  
 tcatggtctt cggagaaatc 260

<210> 2167  
 <211> 266  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2167

aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt tctgctcttc 60  
 ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt tacctcagag 120  
 tgggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac 180  
 gcttgccctg agcaggaccc agacagcaa gttggctgcg aaacatgcac caaaaccaac 240  
 ttggtcatgg tcttcggaga aatcac 266

<210> 2168  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2168

agnogntgca cgcgtacgta agctcggaat tcngetcgag gccaggcaag cccactcaa 60  
 ccaccacacc ttctccttcg ttcacgctac ccttttctgc ttcttcttct acctttcaag 120  
 ttttaaaagt ataaagatgg cagagacatt cctattttacc tcagagtcgg tgaacgaggg 180  
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240  
 ggaccagac agcaaagttg cctgcgaaac atgcaccana accaacttgg tcatggtctt 300  
 cggagaaatc acg 313

<210> 2169  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max

<400> 2169

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gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120  
cttcngcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180  
gtcacccega caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgcccttgaac 240  
aggacctga cagcaagggtt gcctgtgaga catgcaccaa gaccaacatg 290

<210> 2170

<211> 261

<212> nucleic acid

<213> Glycine max

<400> 2170

gtcgcangca cgcgtacgta agctcggaaat tcggctcgag ctcgagccga attcggctcg 60  
nngagaaaac acgaccangg ccaaanttga ctacgagaag anngtgcttg acacctgcag 120  
gancatcggc ntcgtcncaa atgatgtggg actggangcc gacaactgca aggtcctcgt 180  
caacatngag cagcanagcc ctganattgc tcagggngta cncggccacc ttaccaaaaa 240  
acctgaagaa attggtgcng g 261

<210> 2171

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 2171

gtgcattcgt acgtaagctc ggaattcngc tcgaggccag gcaagcccca ctcaaccacc 60  
acacctctcc tcgttcacgc tacccttttc tgctcttctt ctacctttca agtttttaaa 120  
gtataaagat ggcagagacn ttcctattta cctcagagtc ggtgaacgag ggacaccctg 180  
acaagctctg cgaccaaadc tccgatgctg tctcgcagc ttgcctcgag caggaccctg 240  
acagcaaagt tgccctgcga acatgcacca aaaccanctt ggtcatggtc ttcggagaaa 300  
tcacg 305

<210> 2172

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 2172

tcgcangcac gcgtacgtaa gctcggaatt ctntctgagg caagccccac tcaaccacca 60  
cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120  
tataaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 180  
caagctctgc gaccaaatct ccgatgctgt cctcgacgct tgcctcgagc aggaccacaga 240  
cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatgggtct tcggagaaat 300  
caga 304

<210> 2173

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 2173

ngangcacgc gtacgtaagc tcggaattcn gctcgaggca agccccactc aaccaccaca 60  
cctctcctcg ttacgctac ccctttctgc tcttcttcta cctttcaagt tttaaaagta 120  
taaagatggc agagacattc ctattttacct cagagtcggt gaacgaggga caccctgaca 180  
agctctgcga ccaaattctcc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240  
gnaagttgc ctgcgaaaca tgcaccanaa ccaacttggc catggtcttc ggagaaatca 300  
cganca 306

<210> 2174

<211> 283

<212> nucleic acid

<213> Glycine max

<400> 2174

nnncanangc acgcgtacgt aagctcgga ttcggctcga gcggctcgag accactctct 60  
ctgctcttct tctaccttct aagtttttaa agtattaaga tggcagagac attcctatct 120  
accttcagag tcagtgaacg agggacaccc tgacaagctc tgcgacaaa tctccgatgc 180  
tgtctctgac gcttgcttg ancaggaccc agacagcaag gttgcctgcg aaacatgcac 240  
caagaccaac ttgggtcatgg tcttcggaga gatcaccacc aag 283

<210> 2175  
 <211> 321  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2175

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 actttcctat tcacatctga atcagtgaac gangggcacc tgacaagctc tgtgaccaga 180  
 nctccgatgc tgtgctcgat gcatgcttgg agcaggacct gacagcaagg ttgcctgtga 240  
 aacctgcacc aagaccaaca tggatgatgg tttcggagag atcanaacca agggccaacng 300  
 tgannaataa ganatgtgcn t 321

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<210> 2176  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2176  
 aagencact caaccaccac accactctct ctgntctnct tctacctttc aagtttttaa 60  
 agtattaaga tggcanagac attcctatct acctcagagt cagtgaacga nggacaccct 120  
 gacaagctct ggcaccaa atnccgatgct gtcctcgacg cttgccttga acaggaccca 180  
 gacagcaagg ttgctgcgga aacatgcacc agaccacttg gtcatggctct tngaganatc 240  
 accaccaagg ccacgttgac tacgaganga tcgtgcgtga cacctgcaga acatcggctt 300  
 cgtt 304

<210> 2177  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2177  
 cacgcgtacg taagctcgga attcggctcg aggcagactt ancaacagca caaagcgggt 60  
 tactgtcngt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
 aaaatggcnc aagaaacttt cntattcaca tctgaatcag ngaacgagg gcaccctgac 180



aagctctgtg accagatctc cgatgctgtg ctcgntgcat gcttggagca ggaccctgan 240  
 agcaaggttg cctgtgaaac ctggcaccan gaccaacatg gtgatggttt tcggaga 297

<210> 2178  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2178

gtacangcac gcgtacgtaa gctcggaatt cggctcgagg cagacttaac aacagcacia 60  
 agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120  
 agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
 ccctgacaag ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga 240  
 ccctgacagc aagggttgct gtgaaacctg caccaagcca acatgggtgat ggttttcggg 300  
 gagatcacia 310

<210> 2179  
 <211> 278  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2179

cggtacgtn agctcggaat tcggctcgag cttacaaca gcacaaagcg gggttactgtc 60  
 tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa gttaaaatgg 120  
 cccaagaaac tttctattc acatctgaat cagtgaacga ggggcaccct gacaagctct 180  
 gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct gacagcaagg 240  
 ttgcctgtga aacctgcacc aagaccaaca tggatgat 278

<210> 2180  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2180

cgcatgcacg cgtacgtaag ctcggaattc ggctcgagca acagcnaaaa gcgggttact 60  
 gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctccttgcca gaagttaaaa 120

tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac cctgacaagc 180  
tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240  
aggttgcttg tgaacactgc accaagacca acatggtgat g 281

<210> 2181  
<211> 305  
<212> nucleic acid  
<213> Glycine max  
<400> 2181

gnnnangcac gcgtacgtna gctcggaatt cggctcgagg gccaggcaag ccccaactcaa 60  
ccaccacacc tctcctcgnn cacgctgacc cctnctgct cttctttctac ctttcaagtt 120  
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
accctgacaa gctctgcgac caaatctccg atgcngtctt cgacgcttgc ctgcagcagg 240  
accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300  
gagaa 305

<210> 2182  
<211> 277  
<212> nucleic acid  
<213> Glycine max  
<400> 2182

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag cacaaagcgg gtcactgtct 60  
gttcaagcta ccatctctct ctctctttct tagtgcttcc ttgccagaag ttaaaatggc 120  
ccaagaaact ttctatttca catctgaatc agtgaacgag gggcaccctg acaagctctg 180  
tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggaccctg acagcaaggt 240  
tgctgtgaa acctgcacca agaccaacat ggtgatg 277

<210> 2183  
<211> 187  
<212> nucleic acid  
<213> Glycine max  
<400> 2183

tagtgcttcc ttgccagaag ttaaaatggc ccaagaaact ttctatttca catctgaatc 60

agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
 atgcttggag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
 ggtgatg 187

<210> 2184  
 <211> 282  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2184

tcgcangcac gcgtacgtaa gctcggaatt cngctcgagc aacagcacia agcgggttac 60  
 tgtotgttca agctaccatc tctctctctc tttcttagtg cctccttgcc agaagttaaa 120  
 atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 180  
 ctctgtgacc agatctccga tgctgtgctc gatgcatgct tggagcagga ccctgacagc 240  
 aaggttgctt gtgaaacctg caccaagacc aacatggtga tg 282

<210> 2185  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2185

gtcgcangca cgcgtacgta agctcggnat tcggctcgan ctcgagccga attcgggctc 60  
 gantatacaa cagcaciaaag cgggactact gtctgttcaa gactaccatc tctntctctc 120  
 tttcttagtg cctccttgcc agaagttaaa atggcccaan aaactttcct attcacatct 180  
 gaatcngtga acgaggggca ccctgacaag ctctgtgacc agatctccga tgctgtgctc 240  
 gatgcatgct tggagcagga ccctgacagc aaggttgctt gtgaaacctg caccaagacc 300  
 aacatggtga tggtt 315

<210> 2186  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2186

anacgcangc acgcgtacgt aagctcgga ttcngctcga gggcaagccc cactcaacca 60

ccacacctct cctcggtcac gctacccctt tctgctcttc ttctaccttt caagttttaa 120  
aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180  
tgacaagctc tgcgaccaa tctccgatgc tgcctcgcac gcttgccctc agcaggaccc 240  
agacagcaaa gttgctgcg aaacatgnac caaaaccaac ttggtcatgg tcttcggaga 300  
aat 303

<210> 2187  
<211> 297  
<212> nucleic acid  
<213> Glycine max  
<400> 2187

acgtgcgacg cagcgtacg taagctcgga attcngctcg aggccccact caaccaccac 60  
accnctcttc gttcacgcta cccctttctg ctcttcttct acctttcaag ttttaaaagt 120  
ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg acaccctgac 180  
aagctctgcg accaaatctc cgatgctgtc ctgcacgctt gcctcgagca ggaccagac 240  
agcaaagttg cctgcgaaac atgcaccaa accaacttgg tcatggctct cggagaa 297

<210> 2188  
<211> 276  
<212> nucleic acid  
<213> Glycine max  
<400> 2188

cgcntgcacg cgtacgttag ctcggaattc ggctcgaggc acaaagcggg ttactgtctg 60  
ttcaagctac catctctctc tctctttctt agtgccctct tgccagaagt taaaatggcc 120  
caagaaactt tctattcac atctgancca gtgaacgagg ggcaccctga caagctctgt 180  
gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga cagcaagggt 240  
gcctgtgaaa cctgcaccaa gaccaacatg gtgatg 276

<210> 2189  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2189

ngtcgcangc acgcgtacgt aagctcggga attcngctcg aggcaagccc cactcaacca 60  
 ccacacctct cctcggtcac gctacccctt tctgctcttc ttctaccttt caagttttta 120  
 aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180  
 tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgctcgc agcaggaccc 240  
 agacagcaaa gttgctcgcg aaacatgcac caaaaccaac ttggtcatgg tcttcggaga 300

<210> 2190  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2190

gentangtac gcnacgtaa gctcggaatt cggtcgcagc caccacacct ctctcgttc 60  
 acgtacccc ttctngctct tcttctacct ttcaagtttt aaaagtataa agatggcaga 120  
 gacattccta ttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 180  
 aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcttg 240  
 cgaaacatgc accaaaacca acttggtcat ggtcttcgga gaa 283

<210> 2191  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2191

tcgtcgcang cagcgtacg taagctcgga attcggctcg agaggcaagc ccactcaac 60  
 caccacacct ctctcgttc acgtacccc ttctgctct tcttctacct ttcaagtttt 120  
 aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180  
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240  
 ccagacagca aagttgcttg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 300  
 gaa 303

<210> 2192  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max

[illegible]

<210>	2193
<211>	301
<212>	nucleic acid
<213>	Glycine max

agttnnngcac	gcgtacgtaa	gctcgggaatt	cggctcgaga	ggcaagcccc	actcaaccac	60
cacacctctc	ctcgttcacg	ctaccccttt	ctgctcttct	tctacctttc	aagtttttaa	120
agtataaaga	tggcagagac	attcctattt	acctcagagt	cgggtgaacga	gggacaccct	180
gacaagctct	gcgaccaa	atccgatgct	gtcctcgacg	cttgcoctoga	gcaggaccca	240
gacagcaaag	ttgcctgcga	aacatgcacc	aaaaccaact	tggtcatggt	cttcggagaa	300
a						301

<210>	2194
<211>	284
<212>	nucleic acid
<213>	Glycine max

gcatgcacgc	gtacgtaagc	tcggaattcg	gctcgagcca	agccccactc	aaccaccaca	60
ccactctctc	tgtctcttct	ctacctttca	agttttttaa	gtattaagat	ggcagagaca	120
ttcctattta	cctcagagtc	agtgaacgag	ggacaccctg	acaagctctg	cgaccaaato	180
tccgatgctg	tcctcgacgc	ttgccttgaa	caggaccocag	acagcaagggt	tgootgogaa	240
acatgcacca	agaccaaactt	ggtcatggtc	tccgagagat	cacc		284

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<211> 288  
<212> nucleic acid  
<213> Glycine max

<400> 2195

ncacgtcgca ngcacgcnta cgtaagctcg gaattcggct cgagcaacag caciaagcgg 60  
gttactgtct gttcaagcta ccatctctct ctctctttct tagtgctcc ttgccagaag 120  
ttaaaatggc ccaaganact ttcctattca catctgaatc agtgaacgag gggcaccttg 180  
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggaccctg 240  
acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtgatgg 288

<210> 2196  
<211> 292  
<212> nucleic acid  
<213> Glycine max

<400> 2196

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ttcttctngct tgagaaatgg cacaaaaacc tttctattca catctgaatc tgtaaacgan 180  
ggtcaccccg acaagctgtg cgaccagatc tctgatgcag tgctcgatgc gtgccttgaa 240  
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<400> 2197

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aagttttaaa agtataaaga tggcaganac attcctatth acctcagagt cggatgaacga 180  
gggacaccct gacaagctct gcgaccaaht ctccgatgct gtctctgacg cttgcntcga 240  
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cttcggagaa atcacg 316

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 ccctgacaag ctctgcgacc aaatctccga tgctgtcttc gacgcttgcc tcgagcagga 240  
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 agaaa 305

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 ttcttcagct tgagaaatgg cacaagaaac ctttctattc anatctgaat ctgtaaacga 180  
 gggtcacccc gacaagctgt gngaccagat ctctgatgca gtgcccgatg cgtgccttga 240  
 acaggncctt gacancaagg ttgcctgtga gacatgnacc aagaccaana tggatcatgtt 300  
 t 301

<210> 2200  
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 <212> nucleic acid  
 <213> Glycine max  
  
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 gtgcgancga cgcgtacgta agctcggaa ttcggctcga gacttaacaa cagcacaag 60  
 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120  
 aagttaaaat ggccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180



ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcattgcttg gagcaggacc 240  
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<210> 2201  
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<212> nucleic acid  
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gtcgcattgca cgcctacgta agctcggaat tnnctcagag ggccaggcaa gcccactca 60  
accaccacac ctctctctgt tcacgctacc cttttctggc tcttcttcta cttttcaagt 120  
tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 180  
caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240  
gaccagaca gcaaagttgc ctgcgaaaca tgcaccanaa ccaacttggt catggtcttc 300  
ggagaaatc 309

<210> 2202  
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<212> nucleic acid  
<213> Glycine max  
  
<400> 2202

gcagacttaa caacagcaca aagcgggtta ctgtctgttc aagctaccat ctctctctct 60  
nctttcttag tgctctcttg ccagaagtta aaatggccca agaaactttc ctattcacat 120  
ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 180  
tcgatgcattg cttggagcag gaccctgaca gcaaggttgc ctgtgaaacc tgcaccaaga 240  
ccaacatggt 250

<210> 2203  
<211> 295  
<212> nucleic acid  
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ntctctctgt tcacgctacc cttttctgct cttcttctac ctttcaagtt ncaaaagtat 120

aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa 180  
gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 240  
caaagttgcc tgcgaaacat gcaccaaaac caacttggtc atggtcttcg gagag 295

<210> 2204  
<211> 272  
<212> nucleic acid  
<213> Glycine max  
  
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gncgengcac gcgtagttna gctcgggnatt cggctcgagg gcccactca accaccacac 60  
cactctctct gotcttcttc tacctttcaa gtttttaaag tattaagatg gcagagacat 120  
tctatattac ctcagagtca gtgaacgagg gacacctga caagctctgc gaccaaattct 180  
ccgatgctgt cctcgacgct tgccttgaac aggaccaga cagcaagggt gcctgcgaaa 240  
catgcaccaa gaccaacttg gtcattggtct tc 272

<210> 2205  
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<212> nucleic acid  
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<400> 2205

cgtcgcangc acgcgtacgt aagctcggaa ttcggctcga gccaaagccc actcaaccac 60  
cacaccactc tctctgctct tcttctacct ttcaagtttt taaagtatta agatggcaga 120  
gacattccta ttacctcag agtcagtga cgagggacac cctgacaagc tctgcgacca 180  
aatctccgat gctgtctctg acgcttgctt tgaacaggac ccagacagca aggttgcttg 240  
cgaaacatgc accaagacca acttggtcat ggtctt 276

<210> 2206  
<211> 307  
<212> nucleic acid  
<213> Glycine max  
  
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ntontatgca cgcgtacgta agctcggaa ttcggctcgag ggccaggcaa gcccactca 60  
accaccacac ctctctctgt tcacgctacc cttttctgct cttctcttac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240  
 acccagacag caaagttgcc tgcgaaacat gcacccaaac caacttggtc atggtcttcg 300  
 gagaaat 307

<210> 2207  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2207

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 tgcaagctct gottccagcg agtggttcttt ctctgtttca acaccttaat ttgcacacgc 120  
 tgctttctca gottgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180  
 cgaggggtcac cccgacaagc tgtgcgacca gatctctgat gcagtgcctg atgcgtgcct 240  
 gaacaggacc ctgacagcaa ggttgctgtg gagacatgca ccaagaccaa catggtcagg 300  
 tcttgagag a 311

<210> 2208  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
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 tagtcatggt ggtgtttttg gctgtgaatt tgctcatatg tgctaattat gtgttcttgt 120  
 ttgatgttac tctacagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 180  
 agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctnccgatg 240  
 catgcttggg gcaggaccct gacagcaang ttgcctgtga aacctgcacc aagaccaaca 300  
 tggatgatgt 310

<210> 2209  
 <211> 338  
 <212> nucleic acid  
 <213> Glycine max

<400> 2209

tgcgatgcac gcgtacgtaa gctcgggaatt cnnctcgagg caagccccac tcaaccacca 60  
cacctctect cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120  
tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacaccctga 180  
caagctctgc gaccaaactc cgatgctgtc ctcgacgctt gcctcgagca ggaccagac 240  
agcaaagttg cctgcgaaac atgcaccaa accaacttgg tcatggtctt cggagaaatc 300  
acgaccaggc caagttgatt acgagaagta gtgcgtga 338

<210> 2210

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2210

antencangc acgcgtacgt aagctcggaa ttcggctcga gaacagcaca aagcgggtta 60  
ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc cagaagttaa 120  
aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc accctgacaa 180  
gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttgganccag gaccctgaca 240  
gcaaggttgc ctgtgaaacc tgcaccaaga ccaacatggt gatggttt 288

<210> 2211

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2211

gtcgcatgca cgcgtacgta agctcggaa tccggtcgag ggccaggcaa gccccactca 60  
accaccacac ctctcctcgt tcacgctacc ctttctgctc ttcttctacc tttcaagttt 120  
taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180  
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240  
ccagacagca aagttgcctg cgaaacatgc accaaaacca acttggtcat ggtcttcgga 300  
gaaatcacga c 311

<210> 2212

<211> 328  
 <212> nucleic acid  
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 angtcctcang cacgcgtacg taagctcgga attcagctcg agcngctcga gcacacctct 60  
 cctcggttcac gctacccctt tctgctctnc ttctaccttt caagtttttna angtnntaaag 120  
 gtggcagaga cattcctatt tacctcagag tcgntgaacg agggacaccc tgnnaagctc 180  
 tgcgaccaa tctccgatgc tgtcctcgac gcttgccctcg agcaggaccc agacagnaaa 240  
 gttgcntgcg aaacatncac caaaaccaat tggatcatggt cttcggagaa atcacgacca 300  
 aggccaaagt tgatacgaga agatatgc 328

<210> 2213  
 <211> 309  
 <212> nucleic acid  
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 <400> 2213  
  
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 agtataaaga tggcagagac attcctatct acctcagagt cggatgaacga gggacaccct 180  
 gacaagctct ggcaccaaatt ctccgatgct gtccctcgacg cttgcctcga gcaggaccca 240  
 gacagcaaag ttgcctgcga aacatgcacc aaaaccaatt ggtcatgggtc ttcggagaaa 300  
 tcacgacca 309

<210> 2214  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2214  
  
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 tcgacttcag acctggaatg atcaccatta acttgacact taagaggggt ggtcataggt 120  
 tcctcaagac agctgcttat ggacactttg gaaggatga tgcagacttc acctgggaag 180  
 ttgtgaagcc actcaagtca gagaagcctc aagcttaaga gtgttggtta gttaatcact 240

cccttcagtg gatgtcttgc tgggtgtgga tgaataattt gcgtgtttca tgactacta 299

<210> 2215  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max

<400> 2215

ngtgcangc acgcgtacgt aagctcggaa ttcggctcga ggccaggcaa gcccactca 60

accaccacac ctctcctcgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180

accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240

accagacag caaagttgcc tgcgaaacat gcacccaaac caacttggtc atggtct 297

CCCTTCAGTG  
 GATGTCTTGC  
 TGGGTGTGGA  
 TGAATAATTT  
 GCGTGTTTCA  
 TGACTACTA  
 NGTGCANGC  
 ACGCGTACGT  
 AAGCTCGGAA  
 TTCGGCTCGA  
 GGCCAGGCAA  
 GCCCCTCA  
 ACCACCACAC  
 CTCTCCTCGT  
 TCACGCTACC  
 CTTTTCTGCT  
 CTTCTTTCTAC  
 CTTTCAAGTT  
 TTAAGAGTAT  
 AAAGATGGCA  
 GAGACATTCC  
 TATTACCTC  
 AGAGTCGGTG  
 AACGAGGGAC  
 ACCCTGACAA  
 GCTCTGCGAC  
 CAAATCTCCG  
 ATGCTGTCCT  
 CGACGCTTGC  
 CTCGAGCAGG  
 ACCAGACAG  
 CAAAGTTGCC  
 TCGGAAACAT  
 GCACCCAAAC  
 CAACTTGGTC  
 ATGGTCT

<210> 2216  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max

<400> 2216

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accaccacac ctctcctcgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120

ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180

accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240

accagacag caaagttgcc tgcgaaacat gcaccannac caacttggtc atggtctt 298

<210> 2217  
 <211> 284  
 <212> nucleic acid  
 <213> Glycine max

<400> 2217

tcgcangcac gcgtacgtaa gctcgggaatt cggctcgagc ttaacaacag cacaagcgg 60

gttactgtct gttcaagcta ccattctctct ctctctttct tagtgctcc ttgccagaag 120

ttaaaatggc ccaagaaact ttctattca catctgaatc agtgaacgag gggcaccctg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttgag caggaccctg 240

acagcaaggt tgctgtgaa acctgcacca agaccaacat ggtg 284

<210> 2218  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2218

ttgcatgcan gcgtacgtaa nctcggaat tcggctcgag nggagttagg ttctgcacgc 60  
 tctncttcca gcgagtgttc tttctcgtt tcaacacctt aatttgcang acgtgcttn 120  
 tnaaganttg agaaatggca caagaaacct ttctattcac atctgaatct ntaaagcagg 180  
 gtcaccccgga naagctgtgc gancagatct ctnatgcagt gctcgatgcg tgccctgaac 240  
 aggaacctga cagcaaggtt gcctgtgaga catgcaccaa gaccaacatg gcatggtc 298

2218  
298  
nucleic acid  
Glycine max  
2218

<210> 2219  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2219

gtngnnggc nngnacggta cgtgagctcg gaattcggct cgagcaggca agccccactc 60  
 aaccaccaca cctctctcgc ttcacgtac cctttctgc tcttcttcta cctttcaagt 120  
 tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggg gaacgaggga 180  
 caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240  
 gaccagaca gcaaagttgc ctgcgaaaca tgcagcaaaa ccaacttggg catggtcttc 300  
 ggn 303

<210> 2220  
 <211> 301  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2220

ttacatgcac acgtacgtaa gctcggaatt cngctcgaga ggcaagcncc actcaaccac 60  
 cacacctctc ctggttcacg ctaccccttt ctgctcttct totacctttc nagttttaaa 120  
 agtataaaga tggcagagac attcctatct acctcagagt cgggtgaacga gggacaccct 180

gacaagctct gcgaccaa at ctccgatgct gtcctcgacg cttgcctcga gcaggaccca 240  
gatagcaaag ttgcctgcna aacatgcacc aaaaccaact tggatcatggt cttcggagaa 300  
a 301

<210> 2221  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
<400> 2221

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga ggggccaggc aagccccact 60  
caaccaccac acctctcttc gttcacgcta cccctttctg ctcttcttct acctttcaag 120  
ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240  
ggaccagac agcaaagttg cctgcgaaac atgcaccaga accaacttgg tcatggtctt 300  
cgga 304

<210> 2222  
<211> 311  
<212> nucleic acid  
<213> Glycine max  
<400> 2222

tancgcatgc acgcgngtaa nntnnnaatc ggnattcggc tcgagtttga ggccaggcaa 60  
gccccactca accaccacac ctctcctcgt tcacgctacc cctttctgct cttcttctac 120  
ctttcaagtt ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg 180  
aacgagggac accctgacaa gctctgcgat caaatctccg atgctgtcct cgacgcttgc 240  
ctcgagcagg acccagacag caaagttgcc tgcgaaacat gcacaaaac caacttggtc 300  
atggtcttcg g 311

<210> 2223  
<211> 284  
<212> nucleic acid  
<213> Glycine max  
<400> 2223



annctaattgc acgcgtacgt aagctcggaa ttcggctcga gaacaacagc acaaagcggg 60  
 ttactgtctg ttcaagctac catctctctc tctctttctt agtgccctct tgccagaagt 120  
 taaaatggcc caagaaactt tcctattcac atctgaatca gtgaacgagg ggcaccctga 180  
 caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttggagc aggaccctga 240  
 cagcaagggtt gcctgtgaaa cctgcaccaa gaccaacatg gtga 284

<210> 2224  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2224

nncnaannnn cgcattgcacg cgtacgtaag ctccgaattc ggctcgaggc agacttaaca 60  
 acagcacaaa gcgggttact gtctgttcaa gctaccatct ctctctctct ttcttagtgc 120  
 ctcttgcca gaagttaaaa tggcccaaga aactttccta ttcacatctg aatcagtga 180  
 cgagggggcac cctgacaagc tctgtgacca gatctccgat gctgtgctcg atgcatgctt 240  
 ggagcaggac cctgacagca aggttgcttg tgaaacctgc accaagacca acatggtga 299

<210> 2225  
 <211> 324  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2225

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gatttgaggc caggcaagcc 60  
 ccaactcaacc accacacntc tncncgttca cgctaccctt ttctgnctct tcttgetncc 120  
 tttcaagttt taaaagtata aagatggcag agacattcct atttacctca gagtcgggtga 180  
 acgaggggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 240  
 tcgagcagga ccagacagc aaagttgcct gcganacatg caccaaaacc aacttggtca 300  
 tggctctngga gaaatcacga ccaa 324

<210> 2226  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max

<400> 2226

ntcgcangca cgcgtacgta agctcggaat tcggctcgag ggccaggcaa gcccactca 60  
 acnaccacac ctctctctgt tcacgctacc cttttctgct cttcttctac ctttcaagtt 120  
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
 accctgacaa gctctgagac caaatctccg atgctgtcct cgcgcgttgc ctcgagcagg 240  
 acccagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300  
 gaga 304

<210> 2227

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2227

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag caggcaagcc cactcaacc 60  
 accacacctc tctctgttca cgtacccctt ttctgtctct cttctacctt tcaagtttta 120  
 aaagtataaa gatggcagag acatttctat ttacctcaga gtcggtgaac gagggacacc 180  
 ctgacaagct ctgcgaccaa atctccgatg ctgtctctga cgttgccctc gagcaggacc 240  
 cagacagcaa agttgcttgc gaaacatgca gcaaaaccaa cttggtcatg gtcttcggag 300

<210> 2228

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2228

ngtcgcangc acgcgtacgt aagctcgga ttcggctcga gggccaggca agccccactc 60  
 aaccaccaca cctctctctg ttacgctac ccttttctgc tcttcttcta ctttcaagt 120  
 tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg 180  
 caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag 240  
 gaccagaca gnaagttgc ctgcgaaaca tgcacaaaaa ccaacttggt catggtcttc 300  
 gg 302

<210> 2229

<211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2229  
  
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 accacacctc tctctgttca cgtaccctt ttctgctctt cttctacctt tcaagtttta 120  
 aaagtataaa gatggcagag acatttcctat ttacctcaga gtcggtgaac gagggacacc 180  
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240  
 cagacagcaa agttgcctgc gaaacatgca ncaaaaccaa cttggtcatg gtcttcgg 298

<210> 2230  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2230  
  
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 ccaccacacc tctctcggtt cagctaccc ctttctgctc ttcttctacc tttcaagttt 120  
 taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180  
 ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
 ccagacagc aaagttgcct gcgaaacatg caccaaaacc aacttggtca tgggtcttc 298

<210> 2231  
 <211> 269  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2231  
  
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 tttctnctct tcttctacct ttcangtttt aaaagtataa agatggcaga gacattccta 120  
 tttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat 180  
 gctgtcctcg acgcttgctt cgagcaggac ccagacagca aagttgcctg cgaaacatgc 240  
 accaaaacca acttggtcat ggtcttcgg 269

<210> 2232

<211> 290  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2232  
  
 ncaattaana gtcgcangca cgcgtacgta agctcggaat tcggctcgag gttaggttct 60  
 gcacgctctg ctccagcga gtgttctttc ttcgtttcaa caccttaatt tgcacacgct 120  
 gcttcttcag cttgagaaat ggcacaagaa acctttctat tcacatctga atctgtaaac 180  
 gagggtcacc ccgacaagct gtgcgaccag atctctgatg cagtgcctga tgcgtgcctt 240  
 gaacaggacc ctgacagcaa ggttgccctgt gagacatgca ccaagaccaa 290

<210> 2233  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2233  
  
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 accaccacac ctctctctgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120  
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240  
 accagacag caaagttgcc tgcgaaacat gcaccanaac caacttggtc atggtcttcg 300  
 gagaaa 306

<210> 2234  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2234  
  
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 accaccacac acgctctctg tncacgctac ccttttctgg ctcttcttct accnttcaag 120  
 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca 240  
 ggaccagaca gcaaagttgc ctgcgaaaca tgcacaaaaa ccaacttggc catggtcttc 300

ggagaaatca c

311

<210> 2235  
<211> 289  
<212> nucleic acid  
<213> Glycine max

<400> 2235

natcgcatgc acgcgtacgt nagctcggaa ttcggctcga gcaacagcac aaagcggggtt 60  
actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg ccagaagtta 120  
aatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg caccctganc 180  
aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca ggaccctgac 240  
agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatggttt 289

<210> 2236  
<211> 260  
<212> nucleic acid  
<213> Glycine max

<400> 2236

agcagactta acaacagcac aaagcggggtt actgtctgtt caagctacca nnnnnnnnnn 60  
nnnnnnntag tgcctccttg ccagaagtta aatggccca agaaactttc ctattcacat 120  
ctgaatcagt gaacgagggg caccctncac aagctctgtg accagatctc cgatgctgtg 180  
ctcgatgcat gcttggagca ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag 240  
accaacaatgg tgatggttt 260

<210> 2237  
<211> 298  
<212> nucleic acid  
<213> Glycine max

<400> 2237

ntcanntacg cgtangtanc actgcgtacn tnagctcgga attcggctcg agcagcacia 60  
agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120  
agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180  
ccctgtacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240

accctgacag caaggttgcc tgtgaaacct gcaccaagac caacatgggtg atgggtttt 298

<210> 2238  
<211> 301  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2238

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acgctcngcg tnnagngagt gatntttctt cgtntcanca cntnaaattg cancacgctg 120  
cttcttcngc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180  
anggtcaccg cgacaagctg tgtgaccaga tctctgatgc antgctcgat gcgngccttg 240  
aacaggaccc tgacagcaag ttgctgtga gacatgcacc atgaccaaca tggtcaggtc 300  
n 301

<210> 2239  
<211> 309  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2239

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ccacacctct cctcgttcac gctaccctt tctgtactt cttctacctt tcaagtttta 120  
aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180  
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgcttc gagcaggacc 240  
cagacagcaa agttgcctgc gaacatgcac caaaaccaac ttggtcattg tcttcggaga 300  
aatcacgac 309

<210> 2240  
<211> 293  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2240

ngtcgcangc acgcgtacgt aagctcgga ttcggctcga gtaacaaca gcacaaagcg 60  
ggttactgtc tggtcangct accatctctc tctctctttc ttagtgcttc cttgccagaa 120

gttaaaatgg cccaagaaac ttctctattc acatctgaat cagtgaacga ggggaccctg 180  
acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgcttggag caggncctg 240  
acagcaaggt tgcctgtgaa acctgcacca agaccaacat ggtgatgggt ttc 293

<210> 2241  
<211> 279  
<212> nucleic acid  
<213> Glycine max

<400> 2241

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tgtgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag aagttaaaat 120  
ggcccaagaa actttctctat tcacatctga atcagtgaac gaggggcacc ctgacaagct 180  
ctgtgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc ctgacagcaa 240  
ggttgctctg gaacctgcac caagaccaac atgggtgatg 279

<210> 2242  
<211> 181  
<212> nucleic acid  
<213> Glycine max

<400> 2242

tagtgctcc ttgccagaag ttaaaatggc ccaagaaact ttctatttca catctgaatc 60  
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
atgcttggag caggaccctg acagcaaggt tgcctgtgaa acctgcacca agaccaacat 180  
g 181

<210> 2243  
<211> 289  
<212> nucleic acid  
<213> Glycine max

<400> 2243

acgtcgcang cacgcgtacg taagctcgga attcggctcg aggcagactt aacaacagca 60  
caaagcgggt tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt 120  
gccagaagtt aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg 180

gcaccctgac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca 240  
ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaacaatg 289

<210> 2244  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2244

nnocgancga cgcgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60  
aagcgggtta ctgtctgttc aagctaccat ctctctctct ctttcttagt gcctccttgc 120  
cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180  
accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
accctgacag caagggttgc tgtgaaacct gcaccaagac caacatg 287

<210> 2245  
<211> 310  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2245

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caaccancac acatctnctc gttcaagcta cccctttgtn cncncttct aantttcaag 120  
ttttaaaagt atacagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
acacntgac aagctctgcg accaaatctc cgatgctgtc ctcgacgott gcntcgagca 240  
ggaccagac agcaangttg cctgcgaaac atgcacnga accaacttgg tcatggtctt 300  
cggagaaatc 310

<210> 2246  
<211> 284  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2246

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tcacgctacc cctttctgct cttcttctac ctttcaagtt tnaaangtnt aaagatggca 120



gagacatncc tatttacctc agagtcggtg aacgagggac acccngacaa gctctgcgac 180  
 caaanctccg atgcngtcct cgacgcttgc ctcgagcagg acccagacag caaagntgcc 240  
 tgcgaaacan gcacaaaaac caacttggtc atggtcttcg gaga 284

<210> 2247  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2247

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 aaccaccaca ctggctcctc gttnacgcta cccctttctn cctctttctc tacctttcaa 120  
 gttttaaaag tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg 180  
 gacaccctga caagctctgc gaccaaactc ccgatgctgt cctcgacgct tgccctgagc 240  
 aggaccaga cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatggtc 299

<210> 2248  
 <211> 182  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2248

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 agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
 atgcttgag caggaccctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180  
 gg 182

<210> 2249  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2249

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 ancnggtta cttnctgtnc aagctancca tctctctctc tctttcttag tgctccttg 120  
 ccagaagtta aaatggccca agaaactttc ctannacat ctgaatcagt gaancgagg 180

gcacctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcncg nttggagcag 240  
gaccctgaca gcaaggttgc ctgtgaaacc ngcaccaaga ccaacatggt gatgggttttc 300  
ggaganntca caa 313

<210> 2250  
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<212> nucleic acid  
<213> Glycine max  
<400> 2250

gcacgcgtac gtaagctcgg aattcggctc gagcaggcaa gcccactca accaccacac 60  
ctctctctgt tcacgctacc cttttctgct cttctttctac ctttcaagtt ttaaaagtat 120  
aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac accctgacaa 180  
gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag 240  
caaagttgcc tgcgaaacat gcancaaaac caacttggtc atggtcttc 289

<210> 2251  
<211> 264  
<212> nucleic acid  
<213> Glycine max  
<400> 2251

atttgaggcc aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt 60  
tctgctcttc ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt 120  
tacctcagag tcggtgaacg agggacaccc tgacaagctc tgcgaccaaa tctccgatgc 180  
tgtcctcgac gcttgctcgc agcaggaccc agacagcaaa gttgcctgcg aaacatgcac 240  
caaaaccanc ttggtcatgg tctt 264

<210> 2252  
<211> 315  
<212> nucleic acid  
<213> Glycine max  
<400> 2252

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acacctctcc tcgttcacgc tacccttttc tgctctttct ctacctttca agtttttaaaa 120

gtataaagat ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg 180  
aaagctctgc gaccaaattct ccgatgctgt cctcgacgct tgcctcgagc aggaccaga 240  
cagcaaagtt gcttgcgaaa catgcacnaa aaccaattgg tcatgggtctt cggagaaatc 300  
acgaccaagg ccaag 315

<210> 2253  
<211> 191  
<212> nucleic acid  
<213> Glycine max  
<400> 2253

tagtgctcc ttgccagaag ttaaaatggc ccaagaaact ttctatttca catctgaatc 60  
agtgaacgag gggcaccctg acaagcttgt gaccagatct ccgatgctgt gctcgatgca 120  
tgcttggagc aggaccctga cagcaaggtt gctgtgaaa cctgcaccaa gaccaacatg 180  
gtgatggttt t 191

<210> 2254  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
<400> 2254

gttgcangca cgcgtacgta agctcggaat tcggctcgag agcagactta acaacagcac 60  
aaagcgggtt actgtctgtt caagctnca nctctctctc tctttcttag tgctccttg 120  
ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacganggg 180  
caccctgtgc aagctctgtg accagatctc cgatgctgtg ctcgatgcat gattggagca 240  
ggaccctgac agcaagggtg nctgtgaaac ctgcaccaag ancaacatgg tgatggtttt 300  
cgga 304

<210> 2255  
<211> 317  
<212> nucleic acid  
<213> Glycine max  
<400> 2255

tgcgatgcnc gcgtacgtna gctcgggnatt cggctcgagc anaangcngg ttactgtctg 60

ttcaagctac catctctctc tctctttctt antgcctcct tgccagangt taaaatggcn 120  
caagaanctt tcttattcac atctgaatnn gtgaacgagg ggcaccctga acaanctctg 180  
tgancagatc tccgatgctg tgctcgntgc atncttggag caggaccctg acagcnaggt 240  
tnctgtgna acntgcacca agnccancat ggngatgggt ttcggagann tcacaaccan 300  
ggccaacgtg gactatg 317

<210> 2256  
<211> 235  
<212> nucleic acid  
<213> Glycine max  
<400> 2256

cngcacgctg acgtaagctc ggaattcggc tcgagggaga ttggtgctgg tgaccaaggt 60  
catatgttgc gctatgnnct gacgangntc ccgagctcat gcccntgagc catgtccttg 120  
ccacgaagct cgggtgtcaag ctcanagagg ttcggaanaa cgggacatgc ccttggtgta 180  
ganctgntgg caagaccnag gtcantgttg nnnactacaa tggcaagggn gccat 235

<210> 2257  
<211> 319  
<212> nucleic acid  
<213> Glycine max  
<400> 2257

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ctctgcttcc agcgagtgtt ctttcttcgt ttcaacaacc ttaatttgca cacgctgctt 120  
cttcagcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180  
gtcaccocga caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccttgaac 240  
aggaccctga cagcaagttg cctgtgagac atgcaccaag accaactggt caggtctttg 300  
gagagatcac aaccagggc 319

<210> 2258  
<211> 306  
<212> nucleic acid  
<213> Glycine max  
<400> 2258

gtcgcacgca cgcgtacgta agctcggaat tcggctcgag caggcaagcc cactcaacc 60  
 accacacctc tctcgttca cgtacccct ttctgctctt cttctacctt tcaagtttta 120  
 aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180  
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240  
 cagacagcaa agttgcctgc gaaacatgca acaaaaccaa ttggtcatgg tcttcggaga 300  
 aatcac 306

<210> 2259  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 2259  
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 cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tcttgccag 120  
 aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggacacc 180  
 ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240  
 ctgacagcaa ggttgccctgt gaaacctgca accaagacca acatggtgat ggnttncgga 300

<210> 2260  
 <211> 330  
 <212> nucleic acid  
 <213> Glycine max

<400> 2260  
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 cactcaacca ccacacctct cctcgttcac gctacccctt aactgcttct tcttctacct 120  
 ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa 180  
 cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct 240  
 cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca cttggtcatg 300  
 gtcttcggag aaatcagacc aaggccaagt 330

<210> 2261  
 <211> 180  
 <212> nucleic acid

<213> Glycine max

<400> 2261

tagtgccctcc ttgccagaag ttaaaatggc ccaagaaact ntcctattca catctggatc 60  
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcnntgc 120  
atgcttggag caggancctg acagcaaggt tgctgtgaa acctgcacca agaccaacat 180

<210> 2262

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2262

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag caagccccac tcaaccacca 60  
cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120  
tataaagatg gcagagacat tcctatttac ctacagatcg gtgaacgagg gacaccctga 180  
caagctctgc gaccaaactc ccgatgctgt cctcgacgct tgctcgagc aggaccaga 240  
cagcaaagtt gctgcgaaa catgcaccaa aaccaacttg gtcatg 286

<210> 2263

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2263

gtcgcangca cgcgtacgta agctcggaat tcggctcgag tttgaggcca ggcaagcccc 60  
actcaaccac cacacctctc ctcgttcacg ctaccctttt ctgctcttct tctacctttc 120  
aagttttaaa agtataaaga tggcagagac attcctatth acctcagagt cggatgaacga 180  
gggacaccct gacaagctct gcgaccaaact ctccgatgct gtctctgacg cttgcctcga 240  
gcaggacca gacagcaaag ttgctgcga aacatgcacc aaaaccnact tggatcatggt 300

<210> 2264

<211> 332

<212> nucleic acid

<213> Glycine max

<400> 2264

cgcangcacg cgtacgtaag ctcggaattc ggctcgagcg acttaacaac agcacaaagc 60  
 gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct cnttgccaga 120  
 agttaaaatg gccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc 180  
 tgacaagctc tgtgaccaga tctccgatgc tgtgctcgnt gcatgcttgg agcaggaccc 240  
 tgacagcaag gttgcctgtg aaacctgcac caagaccaac atgtgatggg ttccggagagn 300  
 tcacaaccan gcaacgtgga ctatgagagg tt 332

<210> 2265  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max

<400> 2265  
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 gtctgttcaa gctaccatct ctctctctct ttcttagtgc ctcccttgcca gaagttaaaa 120  
 tggoccaaaga aactttccta ttcacatctg aatcagtga cggaggggcac cctgacaagc 180  
 tctgtgacca gatctccgat gctgtgctcg atgcatgctt ggagcaggac cctgacagca 240  
 aggttgcttg tgaaacctgc accaagacca acat 274

<210> 2266  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 2266  
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 taacaacagc acaaagcggg ttactgtctg ttcaagctac catctctctc tctctttctt 120  
 agtgctcctt tgccagaagt taaaatggcc caagaaactt tcctattcac atctgaatca 180  
 gtgaacgagg ggcaccctga caagctctgt gaccagatct ccgatgctgt gctcgatgca 240  
 tgcttgagac aggaccctga cagcaagggt gcctgtgaaa cctgcaccaa gaccaacatn 300

<210> 2267  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max

<400> 2267

ngtcgcangc acgcgtacgt aagctcggaa ttcnngctcga ggccccactc aaccaccaca 60  
cntctcctcg ttcacgctac ccttttctgc tcttcttcta cctttcaagt tttaaaagta 120  
taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggga caccctgaca 180  
agctctgcga ccaaattctcc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240  
gnnaagttgc ctgcgaaaca tgcaccaaaa ccaacttggc catggtct 288

<210> 2268

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2268

gttgcangca cgcgtacgta agctcggaa tcggctcgag atttgaggtt tggagcgact 60  
naactaatca ttaatttgca ctgctgttt cagcttcac accctttctt ttgcatcatt 120  
tatatctctt gagaaatggc acaagaaacc tntctattca catctgaatc tgtaaacgag 180  
ggtcaccccg acangctgtg cgancagatc tctgatgcag tacttgatgc gtgccttgaa 240  
caggaccctg acagcaaggt tgccngtgag acatgnacca agaccaacat g 291

<210> 2269

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2269

cacgcgtacg taagctcgga attcggctcg agncaaagga gtgatttgga gtttgagcg 60  
actgaactaa tcattaattt gcactcgtg tttcagcttc atcaccttc ttttgcata 120  
tttatatctc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180  
agggtcaccc cgacangctg tncnaccaga tctctgatgc agtacttgat gcgtgccttg 240  
aacnggaccc tggacagcaa gggtgcctgt gagacatgca ccaagaccaa catggtct 298

<210> 2270

<211> 296

<212> nucleic acid

<213> Glycine max



<400> 2270

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gcagacttaa caacagcaca 60

aagcgggtta ctgtctgttc aagctaccat ctntctcttc tctttcttag tgcctccttg 120

ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180

caccctgaac aagctctgtg accagatctc cgatgctgtg ctcgatgcat gcttggagca 240

ggaccctgac agcaagggtg cctgtgaaac ctgcaccaag accaacaatgg tgatgg 296

<210> 2271

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2271

gtcncatgca cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcacaaaag 60

cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120

aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc 180

ctgacaagct ctgtgaccag atctccgatg ctgtgctcga tgcagtcttg gagcaggacc 240

nncgacagca aggttgcttg tgaaacctgc accaagacca acatggtg 288

<210> 2272

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2272

nagtcgcatg cagcgtacg taagctcgga attcngctcg aggcaagccc cactcaacca 60

ccacacctct cctcgttcac gctaccctt tctgctcttc ttctaccttt caagttttta 120

aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180

tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgctcgc agcaggaccc 240

agacagcaaa gttgcctgcg aaacatggca ccaaaaccaa cttggtcatg gtcttcgga 299

<210> 2273

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2273

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cacacctctc ctcgttcacg ctaccccttt ctgctcttct tctacctttc aagtttttaa 120  
agtataaaga tggcagagac attcctatctt acctcagagt cggatgaacga gggacaccct 180  
gacaagctct gcgaccaaatt ctccgatgct gtccctcgacg cttgcctcga gcaggaccca 240  
gcagacaaag ttgcctgcga aacatgcacc anaaaccaac ttgggtcatgg tcttcggaga 300

<210> 2274

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 2274

acgtcgcang caogcgtagc taagctcggg attcggctcg agggagtttg gagcgactga 60  
actaatcatt aatttgcact cgctgtttca gcttcacac cttctttttg catcatttat 120  
atctcttgag aaatggcaca agaaaccttt ctattcacat ctgaatcgta aacnagggtc 180  
accccgacaa gctgtncnat cagatctctg atgcagtact tgatgcntgc cttgancagg 240  
nccctgacag caaggttgcc tgtgagacnt gcaccaagac caacatgggc atggtct 297

<210> 2275

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2275

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tcctcgttca cgctacccct ttctgctctt cttctacctt tcaagtttta aaagtataaa 120  
gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc ctgacaagct 180  
ctgcgacnna atctccgatg ctgtcctcga cgcttgccct gagcaggacc cagacagcaa 240  
agttgcctgc gaaacatgca ccaaaaccaa ttgggtcatgg tcttcggaga aatcac 296

<210> 2276

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 2276

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accaccacac ctctctctgt tcacgtacc cttttctgct cttttcttac ctttcaagtt 120  
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240  
accagacag caaagttgcc tgcgaaacat gcaccaaacc caattggtca tggctcttcgg 300  
agaaat 306

<210> 2277

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2277

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagg gccaggcaag ccccactcaa 60  
ccaccacacn nctctctggt cagcgtaccc cttttctgctc ttctttctacc tttcaagttt 120  
taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180  
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
cccagacagc aaagttgcct gcgaaacatg caccaaaaacc aacttgg 287

<210> 2278

<211> 206

<212> nucleic acid

<213> Glycine max

<400> 2278

gccacttacc anaaaacctg aagaaattgg tgctgntgac cagggtcaca tgtttggtca 60  
tgccactgat gaaaccctg aattgatgcc attgagccat gttcttgcaa caaaactcgg 120  
tgctcgtctc accgaggttc gcnagaacgg tactggcctt ggctgangct gatggaagac 180  
ccaagtgacc gttgagtata caatga 206

<210> 2279

<211> 265

<212> nucleic acid

<213> Glycine max

<400> 2279

tgcacgcacg cgtacgttag ctcggaattc ggctcgagac agcacaggag cgggttaacng 60  
tctgttcaag ctaccatctc tctctctctt tcntagtgcc tcottgccag aagttaaaat 120  
ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggcacc ctgacaagcn 180  
ctgcgaccag atctccgatg ctgtgctcga tgcattgctg gagcaggacc ctgacagcaa 240  
ggttgccctgt gaaacctgca ccaag 265

<210> 2280

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2280

nanngcncgn gtacgtaagc tcgganttcg gctcgagggc caggcaagcc ccaactcaacc 60  
accacacctc tctcgttca cgctaccctt tngtgcctt cttctacctt tcaagtttta 120  
aaagtataaa gatggcagag acattcctat ttacctcaga gtcgggtgaac gagggacacc 180  
ctgacaagct ctgcgaccaa ntctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240  
cagacagcaa agttgcctgc gaaacatgca ccataaccaa cttgggtcatg g 291

<210> 2281

<211> 330

<212> nucleic acid

<213> Glycine max

<400> 2281

gcacgcgtac gtaagctcgg aattcggctc gagatttgag gccaggcaag cccactcaa 60  
ccaccacacc tctcctcgtt cacgctaacc ctttctgctc ttcttctacc tttcaagttt 120  
tanaagtata aagatggcag agacattcct atttacctca gagtcgggtga acgagggaca 180  
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
cccagacagc aaagttgcct gcgaaacatg gcacaaaac caattgggtca tggctcttcgg 300  
agaaatcaga ccaggccaag ttgattacga 330

<210> 2282

<211> 283

<212> nucleic acid



<211> 208  
 <212> nucleic acid  
 <213> Glycine max

<400> 2285

cangcacgcg tacgtaagct cggaattcgg ctcgagcatt ggtgtccctg agcccttgtc 60  
 agtgtttgtg gacacttatg gaactgggaa gattcctgac aaggagattc tgcaaattgt 120  
 gaaggagaat ttcgacttca gacctggaat gatcaccatt aacttggacc ttaanagggg 180  
 tggatcatagg ttcttcaaga canntgct 208

<210> 2286  
 <211> 270  
 <212> nucleic acid  
 <213> Glycine max

<400> 2286

gcagacttna caacagcaca naggcgggta ctgtctgttc aangctacga tctctgtctc 60  
 tggttatctta gtgcctacct tgccagaagn nannatggcc caatnaantt tccnattcac 120  
 atctgantca ntgaacnatg ggcaccctga naanctctgt gnccagatct ccgatgctgt 180  
 gctcgatgca tgcttggagc aggaccctga nagnaggtt gcntgtnaaa cctgnaccaa 240  
 gaccaacatg gtgatggtn tggagagat 270

<210> 2287  
 <211> 302  
 <212> nucleic acid  
 <213> Glycine max

<400> 2287

nntcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggntttgagg ccaggcaagc 60  
 ccactcaac caccacacnt ctctcgttc acgtacccc tttctggctc ttcttctacc 120  
 tttcaagttt taaaagtata aagatggcag agacattcct atttacctca gagtcgggtga 180  
 acgagggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 240  
 tcgagcagga ccagacagc aaagttgcct gcgaaacatg caccanaacc aacntgggtca 300  
 tg 302

<210> 2288

<211> 195  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2288  
  
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 ttatcaagcc tgtcattctt gagaagtacc ttgatgagaa ncaccatctt ccaccttaac 120  
 ccttctggcc gttttgtcat tgggtggcct catggtgatg ctggtctcac tggaagaaaa 180  
 tcatcattga tacct 195

<210> 2289  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2289  
  
 ngcntgtacg cgtacgtaag ctcggaattc ggctcgaggt caggcaagcc ccaactcaacc 60  
 accacacctc tgectgngtt cangtaccc ctttntgctc ttcttctacc tttgaagttt 120  
 taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acganggaca 180  
 cctgacaag ctctgcganc caaatctccg atgctgtcct cgacgnittgn ctcgagnagg 240  
 acccagacag naaagttgcc tgcgatanat gcaccannac caacttggtc atggtcttcg 300  
 gagaaatcac gacc 314

<210> 2290  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2290  
  
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 ctcaaccacc acacctctcc tcgttcacgc tacccttttc tgctcttctt ctacctttca 120  
 agtttttaaaa gtataaagat ggcagagaca ttcctattta cctcagagtc ggtgaacgag 180  
 ggacaccctg acaagctctg cgaccaaata tccgatgctg tctcgcacgc ttgcctcgag 240  
 caggacccag acagcaaagt tgctgacgaa acatggcacc aaaaccaact tggatcatggt 300  
 ctt 303

<210> 2291  
 <211> 285  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2291  
  
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 ggttactgtc tgttcaagct accatctctc tctctcttct ttagtgctc cttgccagaa 120  
 gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
 gacaagctct gtgaccagat ctccgatgct gtgctcgatg catgcttgga gcaggaccct 240  
 gacagcaagg ttgctgttnn aaacctgcac caagaccaac atggt 285

<210> 2292  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
  
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 aaagcggggtt actgtctgtt caagctacca tctctctctc tctttcttag tgctccttg 120  
 ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180  
 caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag 240  
 gaccctgaca gcaaggttgc ctgtgaaacc tggaccaag accaacatg 289

<210> 2293  
 <211> 343  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2293  
  
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 atacagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg acaccctgac 180  
 aagctctgcg accaaatctc cgatgctgtc ctcgacgctt gcctcgagca ggaacnagac 240  
 agcaaagttg cctgcgaaac atgcacaaa accaattggt catggtcttc ggagaaatca 300



cgacagccan gttgatagag agatatgcgt gacnctgcag aca

343

<210> 2294

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2294

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accacacntc tctcgtttca cgctaccctt tctgctctt cttctacctt tcaagtttta 120

aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180

ctgacancct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240

cagacagcaa agttgcctgc gaaacatgca ccanaaccaa cttggtcat 289

<210> 2295

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2295

gtcgcangca cgcgtacgta nagctcggaa ttcggctcga gntgaggcca ggcaagcccc 60

actcaaccac cacacctctc ctcgttcacg ctaaccctt tctgctctt cttctacctt 120

caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180

agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctg 240

agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaacncac ttggtc 296

<210> 2296

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2296

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ccacacntct cctcgttcac gctaccctt tcttctctt cttctacctt tncaagtttt 120

aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180

cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct cgagcaggac 240

ccagacagca aagttgcctg cgaaacatgc accaaaacca acttgg 286

<210> 2297  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max

<400> 2297

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 accaccacac ctctcctcgt tcacgctacc cttttctgct cttcttctac ctttcaagtt 120  
 ttaaaagtat aaagatggca gngacattcc tatttacctc agagtcggtg aagagggaca 180  
 cctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
 ccagacagc aaagttgcct gcgaaacatg caccagaacc aacttggtca tggctcttcgg 300  
 agaaatcacg accaaggc 318

<210> 2298  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max

<400> 2298

ngnngcacgc gtacgtnagc tcggaattcg gctcgagttt gaggccaggc aagccccact 60  
 caaccaccac acgggctcct cgtnnacgct acccctttct ncctcttctt ctacctttca 120  
 agtttttaaaa gtataaaaat ggacagagaca ttctatttta cctcagagtc ggtgaacgag 180  
 ggacaccctg acaagctctg cgaccaaata tccgatgctg tctcgcagc ttgcctcgag 240  
 caggaccag acagcaaagt tgctgcgaa acatgcacca aaaccaactt 290

<210> 2299  
 <211> 275  
 <212> nucleic acid  
 <213> Glycine max

<400> 2299

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 aaagcgggtt actgtctgtt caagctacca tctctctctc tctttcttag tgcctccttg 120  
 ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt gaacgagggg 180

caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag 240  
gaccctgaca gcaaggttgc ctgtgaaacc tgcac 275

<210> 2300  
<211> 308  
<212> nucleic acid  
<213> Glycine max  
<400> 2300

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cancacangg nnoctcgtn acgtacccc tntctnctc ttcttctacc tttcagngtg 120  
ttaaagtat aaagatggca gagacattcc tatttaccag agtcggtgaa cgagggacac 180  
cctgacaagc totgogacca aatctccgnt gctgtctcgc acgcttgccg cgagcaggac 240  
ccagacagca aagttgcntg cgaaacatgc nccaaaacca acttggnccat ggtcttcgga 300  
gaaatcag 308

<210> 2301  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2301

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aggccccant caaccaccac acctctanct cgttcacgct acccctttct gctcttcttc 120  
taactttcaa gttttaaaag tataaagatg gcagagacat tcctatttac ctcagagtcg 180  
gtgaacgagg gacaccctga caagctctgc gaccaaactc ccgatgctgt cctcgacgct 240  
tgctcgagc aggaccaga cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg 300

<210> 2302  
<211> 295  
<212> nucleic acid  
<213> Glycine max  
<400> 2302

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ccaccacacc tctctcggtt cagctaccc ctttctgctc ttcttctacc tttcaagttt 120

taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggaca 180  
 ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacggcttgc ctcgagcagg 240  
 acccagacag caaagttgcc tgcgaaacat gcancaaaac caacttggtc atggt 295

<210> 2303  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2303

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 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca nacgtgctt 120  
 cttcngcttg agaaatggca caagaaacct ttctattcac atctgaatct gtaaacgagg 180  
 gtcaccccgga caagctgtgc gaccagatct ctgatgcagt gctcgatgcg tgccatgaaca 240  
 ggaccctgac agcaagggtg cctgtgagac atgcaccaag a 281

<210> 2304  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2304

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 getaccatct ctctctctct ttcttagtgc ctcccttgcca gaagttaaaa tggcccaaga 120  
 aactttccta ttcacatctg aatcagtga cgaggggcac cctgagaagt ctgtgaccag 180  
 atctccgatg ctgtgctcga tgcattgntt gagcaggacc ctgacagcaa gggtgcctgt 240  
 naaacctgca ncaagancaa catggtgatg gntttncgga gagatcacia acanngc 297

<210> 2305  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2305

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 ggnacaaagn natcacaagt tcaancnact tntcnnttc ttaangggcg tccntnccag 120

ncgttaaaat ggcccaagaa actttcctat ncacatctga atcagtgaac gangggcacc 180  
ctgacagctc tgtgaccaga tctccgatgc tgtgctcgnt gcntgcntgg agcaggaccc 240  
tgacagcnag gntgcctgtg aaacctgcac caagacnnac atggtgangg ttttcggang 300  
anatcacaac 310

<210> 2306  
<211> 295  
<212> nucleic acid  
<213> Glycine max  
<400> 2306

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ccaccacacc tctcctcggt cactctaccc ctttctgctc ttcttctacc tttcaagttt 120  
taaaagtata aagatggcag agacattcct atttacctca gagtcggtga acgagggacc 180  
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgccct cgagcaggac 240  
ccagacagca aagttgcctg cgaaacatgc accannacca acttggtcat ggtct 295

<210> 2307  
<211> 158  
<212> nucleic acid  
<213> Glycine max  
<400> 2307

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cgaacaggac tcagacagca aggttgccctg cgaaacatgc accaagacca acttggtcat 120  
ggtcttcgga gagacaccac caaggccaac gttgacta 158

<210> 2308  
<211> 302  
<212> nucleic acid  
<213> Glycine max  
<400> 2308

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gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga 120  
agttaaaatg gcccaagaaa ctttcctatt cacatctgaa tcagtgaacg aggggcaccc 180

tgacaagctc tgtgancaga tctccgatgc tgtgctcgat gcatgcttgg agcaggaccc 240  
 tgacagcaag gttgctgtga aacctgcacc aagaccaaca tgtgatgggtt ttcggaanag 300  
 at 302

<210> 2309  
 <211> 295  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2309

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 caccacacct ctctctgttc acgtacnnc tttctgtctt tttctacct ttcaagtttt 120  
 aaaagtataa agatggcaga gacattccta tttacctcag agtcggtgaa cgagggacac 180  
 cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240  
 ccagacagca aagttgcttg cgaaacatgc caccaaaaacc aacttggtca tggtc 295

<210> 2310  
 <211> 271  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2310

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 aagcgggtta ctgtctgttc aagctacat ctctctctt cttcttagt gcctccttgc 120  
 cagaagttaa aatggcccaa gaaactttcc tattcacatc tgaatcagtg aacgaggggc 180  
 accctgacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc ttggagcagg 240  
 accctgacag caaggttgcc tgtgaaacct g 271

<210> 2311  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2311

tnacangcac gcgtacgtaa gctcggaatt cggctcgagg gccaggcaag cccactcaa 60  
 ccaccacacc tctctcgtt cagctaccc cttctgtctc ttctctacc tttcaagttt 120

taaaagtata aagatggcag agacattcct atttacctca gagtcgggtga acgagggaca 180  
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
cccagacagc aaagttgcct gcgaaacatg caccanaacc aattgggtcat ggtttcggag 300  
aatcac 306

<210> 2312  
<211> 308  
<212> nucleic acid  
<213> Glycine max  
<400> 2312

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accaccacac ctctcctcgt tcacgctacc cttttctgct cttctcttac ctttcaagtt 120  
ttaaagtat aaagatggca gagacattcc tatttacctc agagnnggtga acgagggaca 180  
ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
cccagacagc aaagttgcct gcgaaacatg caccanaacc aacttgggtca tggctcttcgg 300  
agaaatca 308

<210> 2313  
<211> 290  
<212> nucleic acid  
<213> Glycine max  
<400> 2313

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cagcaciaaag cgggttactg tctgntcaag ctaccatctc tctctctctt tcttagtgcc 120  
tccttgccag aagttaaaat ggccaanna actttccnat tcacatctga atcagtgaac 180  
gaggggcacc ctgacaagct ctgtgaccag atctccgatg ctgtgctcga ngcatgcttg 240  
gagcaggacc ctgacagcaa ggttgctgtg gaaacctgca ccaagaccaa 290

<210> 2314  
<211> 294  
<212> nucleic acid  
<213> Glycine max  
<400> 2314

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 cacctctcct cgttacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120  
 tataaagatg gcagagacat tcctatttac ctncagagtc ggtgaacgag ggacaccctg 180  
 acaagctctg cgaccaaata tccgatgctg tctcgcacgc ttgcctcgag caggaccctg 240  
 acagcaaagt tgctgcgaa acatgcacca aaaccaactt ggtcatgggc ttcg 294

<210> 2315  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2315

ncgtcgcang cagcggtacg taagctcgga attcggctcg aggatttgag gccaggcaag 60  
 cccactcaa ccaccacacc tctcctcggt cagctaccc ctttctgctc ttcttctacc 120  
 tttnaagttt taaaagtata aagatggcag agacattcct atttacctca ggtcgggtga 180  
 acgaggggaca ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc 240  
 tcgagcagga ccagacagc aaagttgcct gcgaaacatg caccaaaacc aantggt 297

<210> 2316  
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 <212> nucleic acid  
 <213> Glycine max  
 <400> 2316

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 cntccttgct agangttana atggcccang anactttcct attcacatct gantnagtga 120  
 acgaggggca ccctgacaag ctctgtgncc agatctccga tgctgtgctc gatgcatgcn 180  
 tggagcagga ccctgacanc aaggttgctt gtgaaacctg caccaagacc anc 233

<210> 2317  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2317

tcgtgcacg cgtaogtaag ctcggaattc ggctcgaggc caggcaagcc cactcaacc 60



accacacctc tctctgttca cgctaccctt tttctgtctt cttctacctt tcaagtttta 120  
aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180  
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240  
cagacagcaa agttgcctgc gaaacatgca ccaaaaccaa ttggtcat 288

<210> 2318  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
<400> 2318

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tcgagggccc cactcaacca ccacacctct anctcgttca cgctaccacc tttctgtctt 120  
tctttctaaact ttcaagtttt aaaagtataa agatggcaga gacattccta tttacctcag 180  
agtcggtgaa cgagggacac cctgacaagc tctgcgacca aatctccgat gctgtcctcg 240  
acgcttgccct cgagcaggac ccagacagca aagttgcctg cgaaacatgc accaaaacca 300  
attg 304

<210> 2319  
<211> 305  
<212> nucleic acid  
<213> Glycine max  
<400> 2319

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cactcaacca ccacacctct cctcgttcac gctaccctt tctgtcttct ttctaccttt 120  
caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180  
agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgccctcg 240  
agcaggaccc agacagcaaa gttgcctgcg aaacatgcac caaaaccaan tgggtcatggt 300  
cttcg 305

<210> 2320  
<211> 299  
<212> nucleic acid  
<213> Glycine max

<400> 2320

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caccacacct ctctcgttc acgtacccc tttctgctct tttctacct ttcaagtttt 120  
aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180  
cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240  
ccagacagca aagttgcctg cgaaacatgc accaaaacca attggtcatg gtcttcgga 299

<210> 2321

<211> 316

<212> nucleic acid

<213> Glycine max

<400> 2321

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cgttcacgct acccctttct gctcttcttc tacctttcaa gttntaaaag tataaagatg 120  
gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacacctgac aagnctgcga 180  
ccaaactccg atgctgtcct cgacgcttgc ctcgagcagg acccagacag caaagttgcc 240  
tgcgaaacat gcacaaaaac caattgggtca tggctcttcgg agaaatcacg accaggccaa 300  
gttgactacg agaaga 316

<210> 2322

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2322

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tctgcacgct ctgcttcag cgagtgttct ttcttcgttt caacacctta atttgcacac 120  
gctgcttctt cagcttgaga aatggcacia gaaacctttc tattcacatc tgaatctgta 180  
aacgaggggc accccgacaa gctgtgcgac cagatctctg atgagtgtc gatgcgtgcc 240  
ttgaacagga cctgacagc aaggttgctt gtgagacatg caccaaga 288

<210> 2323

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2323

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acanctctcc tcgttcacgc tacccttnc tgcctcttct ctacctttca agtttttaaaa 120  
gtataaagat ggcagagaca ttncatattta cctcagagtc ggtgaacgag ggacaccctg 180  
acaagctctg cgaccaaadc tccgatgntg tccctgaatg naaatcgagc aggaccaga 240  
nagcaaagtt gntgcgana catgcaccaa aaccaacttg gtcatggtct tcggagaaa 299

<210> 2324

<211> 254

<212> nucleic acid

<213> Glycine max

<400> 2324

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tgccctccttg ccagaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt 120  
gaacgagggg caccgcacaa gctctgtgac cagatctccg atgctgtgct cgatgcatgc 180  
ttggagcagg accctgacag caaggttgcc tgtgaaacct ngcaccaaga ccnacatggt 240  
gatggttttc ggag 254

<210> 2325

<211> 227

<212> nucleic acid

<213> Glycine max

<400> 2325

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atttcaggcc tggatgata tccatccaac cttgatctca agaggggtgg aaataacagg 120  
tttttgaaga ctgctgccta tggacacttt ggaagagaag accctgacnt tcacatgggg 180  
aagttggtcc naccctctcn agttgggnaa agccnaacca tttcatc 227

<210> 2326

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 2326

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atttacctac agagtcggtt aacgagggac accctgacaa gctctgacgac caaatcccga 180  
tgctntcttc gagcgttgcc tcgagengga cccagacagc aaagttgcct gcgaaacatg 240  
caccaaaacc aacttggtca tgggtcttcgg aganntcang accaaggcca acgt 294

<210> 2327

<211> 281

<212> nucleic acid

<213> Glycine max

<400> 2327

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anctaccatc tctctctctc tntcttagtg cctcccgtn cagnngtnan aatggcccaa 120  
naaactttcc tattnacatc tgaatcagtg aacgangggc anccganaan ctccgtgacc 180  
agatntccga tgctgtgntc gatgcatgct tggagcagga ccctgacagc aaggttgct 240  
gtganacctg caccaagacc ancatggtgn tggtttncgg a 281

<210> 2328

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 2328

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atggcccann anactttcct atncacatct gaatcantga acgaggggca cnctgacaag 180  
ntctgtgncc agatctccgg tgctgtgctc gatgcatgct tggagcagga ccctgacagc 240  
aagntgcct ggaaacctgc acnaagacca acatggtgat ngntttcgga gann 294

<210> 2329

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2329

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tcaaccacca cacctctcct cgttcacgct acccctttct gctacttctt ctacctttca 120  
agtttttaaaa gtataaagat ggcagagaca ttcctattta cctcagagtc ggtgaacgag 180  
ggacaccctg acangctctg cgaccaaadc tccgatgctg tcctcgacgc ttgcctcgag 240  
caggacccag acagcaaagt tgctgcgaa acatgcacca aaac 284

<210> 2330

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<213> Glycine max

<400> 2330

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tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt gaacgaggga 180  
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<210> 2331

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 2331

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acgaggggca cnctgacaan ntctttgacg agatctccga tgctgtgntc gatgcatgct 240  
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<210> 2332

<211> 262

<212> nucleic acid

<213> Glycine max

<400> 2332

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tcagagtcgg tgaacgaggg acaccctgac aagctntgcg accaaatctc cgatgctgtc 180  
ctcgacgctt gctcgcagca ggaccagac agcaaagttg cnngcgaaac atggcaccaa 240  
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<210> 2333

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2333

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tcggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac 240  
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<400> 2335

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taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca 180  
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<400> 2336

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ggaccagac agcaaagttg cctgcgaaac atgcacaaa acca 284

<210> 2337

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<212> nucleic acid

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<400> 2337

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cttcaactcta cctttcaagt tttaaaagta taaagatggc agagacattc ctatttacct 180  
cagagtcggt gaacgagggg caccctgaca agctctgcga ccaaattctc gatgctgtcc 240  
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2339  
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 acgctacncc tttctgntct tcttctacct ttcaagtttt aaaagtataa agatggcaga 180  
 gacattccta tttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 240  
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 atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca ccctgacaag 180  
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 aaggttgccct gtgaaacct 259

<210> 2341



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 tcaattcaac accttaattt ncacacgctg cttcttcngc ttgaganatg gcacaagaaa 180  
 cntttctatt cacatctgaa tctgtaaagc anggtcacc cgcacaagctg tgcgaccaga 240  
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 agacatgcn 309

<210> 2342  
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 ttaaaatggc ccaanaaact ttctntttca catctgaatc agtgaacgag gggcaccctg 180  
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<210> 2343  
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 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
 acaccctgac aagctctgcg accaaatctc cgatgctgtc ttgcagcgtt gcctcgagca 240  
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 ggccaagntg actacgagaa g 321

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 cttctacctt tcaagtttta aaagtataaa gatggcagag acattcctat ttacctcaga 180  
 gtcggtgaac gagggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga 240  
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 tttgcatcat ttatatctct tgagaaatgg cacaagaaac ctttctattc acatctgaat 180

ctgtaaacga nggtcacccc gacangctgt ncnancagat ctctgatgca gtacttgatg 240  
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 gntccttgg ccanaagtta aaatggccca agaaactttc ctattcacat ctgaatcagt 180  
 gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc tcgatggcat 240  
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 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctc gagcaggacc 240  
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 gtnggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctc 240  
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 ttttaaaagt ataaagatgg cagagacatt cctatttacc tcagagtcgg tgaacgaggg 180  
 acacctgac aagctctgcg accaaatctc cgatgctgtc ctcgncgctt gcctcgagca 240  
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<210> 2357  
 <211> 298  
 <212> nucleic acid  
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 cactcaacca ccacacctct cctcggtcac gctaccctt tctgctcttc ttctaccttt 120  
 caagttttta aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg 180  
 agggacaccc tgacaagctc tgcgacaaa tctccgatgc tgtctctgac gcttgctctg 240

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<210> 2358  
 <211> 288  
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 aacgagggac accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc 240  
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<210> 2359  
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 <212> nucleic acid  
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 tatanagatg gcagagacat tcctatttac ctcagagtcg gtgaacgnng ngacaccctg 180  
 acaagctctg cgaccaaata tccgatgctg tctcgcgcg ttgcctcgag caggacccag 240  
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 tcacgaccaa gccaaattga ctagaagaag atatg 335

<210> 2360  
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 agcagactta acaacatnac aaaccgggtt actgtctgtt canggctacc atctctctct 180

ctctttcttta ggtgnctcct tgccagnang tnnnaatgng gcnaagnaac ttncctattc 240  
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 aataccanca tggatgatggg ttttcggaga gatnccaanc gangccnaan nttgnaacta 420  
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<210> 2361  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max

<400> 2361  
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 gccagaagtt aaaatggccc aagaancctt cctattcaca tctgaatcag tgaacgaggg 180  
 gcacctgac aagctctgtg accagatctc cgatgctgtg tcgatgcatg cttggagcag 240  
 gacctgaca gcaagggtgc ctgtgaaacc tgcaccaaga cca 283

<210> 2362  
 <211> 495  
 <212> nucleic acid  
 <213> Glycine max

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 ccaactcaacc accacacctc nctcgttca cgctaccctt ttctgtctt cttctanctt 180  
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 cganggacac nctgacaagc tctgcgagca aatctccagn tgctgtcctc gacgcttgcc 300  
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 tngtcttcgg anaaatcn cn accaaggcca acnttgactn cnanangata ntgcgttaca 420  
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<400> 2364

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ccctgacagc aaagttggcc tgtgaaacct gcaccangng tgacatgggtg atggtttcg 299

<210> 2366  
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 <212> nucleic acid  
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<400> 2366

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 ctacctttca agtttttaaaa gtataaagat ggcagagaca ttcttattta cctcagagtc 180  
 ggtgaacgag ggacaccctg acaagctctg cgaccaaadc tccgatgctg tctcagacgc 240  
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<210> 2367  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max

<400> 2367

gtcgcngcac gcgtacgtga gctcggaatt cggctcgagg gccaggcaag ccccaactcaa 60  
 ccaccacacc tctcctcggt cacgctaccc ctttctgctc ttcttctacc tttacaagtt 120  
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcgggtg aacgagggac 180  
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240  
 acccagacag caaagttgcc tgcgaaactg caccancacc aattgggtca 289

<210> 2368  
 <211> 302  
 <212> nucleic acid  
 <213> Glycine max

<400> 2368

gtcgcangca cgcgtacgtn agctcggaat tcggctcgag ggccaggcaa gcccactca 60  
 accaccacac ctctcctcgt tcacgctacc ctttctgctc cttctctac ctttcaagtt 120  
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggta acgagggaca 180

ccctgacaag ctctgcgacc aaatctccga tgctgtcctc gacgcttgcc tcgagcagga 240  
 cccagacagc aaagttgcct gcgaaacatg caccataacc aacttggtca tgggtcttcgg 300  
 ag 302

<210> 2369  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2369

cgcatgcacg cgtacgttag ctcggaattc ggctcgagtt tgaggccagg caagcnccac 60  
 tcaaccacca cacctctcct cgttcacgct acccctttct gctacttact tctacctttc 120  
 aagtttttaa agtataaaga tggcagagac attcctattt acctannagc cgggtgaacga 180  
 gggacaccct gacaagctct gcgaccaaatt ctccgatgct gtcctcgacg cttgctctga 240  
 gcaggaccca gacagcaaag ttgcctgcga aacatgcacc aaaacnaa 288

<210> 2370  
 <211> 292  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2370

cgcangcacg cgtacgtaag ctcggaattc ngctcgagca agccccactc aaccaccaca 60  
 cctctctctg ttcacgctac cctttctctg tcttcttcta cctttcaagt tttaaaagta 120  
 taaagatggc agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca 180  
 agctctgcga ccaaattctc gatgctgtcc tcgacgcttg cctcgagcag gaccagaca 240  
 gcaaagttgc ctgcgaaact gcacaaaaac caatgggtca ngntctnaga aa 292

<210> 2371  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2371

gtcgcngcac gcgtacgtaa gctcgggnatt cngctcgagg caagccccac tcaaccacca 60  
 cacctctacc tacgttcacg ctaccccttt ctgctcttct tctaccttta caagttttaa 120

aagtataaag atggcagaga cattcctatt tacctacaga gtcggtgaac gagggacacc 180  
 ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgccctc gagcaggacc 240  
 cagacagcaa agttgcctgc gaaacatgca ccaaaaccaa cttggtca 288

<210> 2372  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2372

acgtcgctgc acgcgtacgt aagctcggaa ttcggtcga ggtgatttgg agtttggagc 60  
 gactgaacta ntcattaatt tgcacttcgc tgtttcagct tcatcaccct tcttttgcac 120  
 catttatatc tcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180  
 ccanggtcac cccgacaagc tgtgcgacca gatctctgat gcagtaacttg atgcntgcct 240  
 gnnccaggacc ctggacagcn aggtancctg tnagacatgc accaagacca acatggtca 299

<210> 2373  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2373

cgcatgcacg cgtacgtnag ctcggaattc ggctcgaggg caagccccac tcaaccacca 60  
 cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120  
 tataaagatg gcagagacat tcctatttac ctccagantcg gtgancgagg gncaccctga 180  
 caagctctgc gaccaaattct ccgatgctgt cctcgacgct ngcctcganc aggacccana 240  
 cagcnaagtt gcctncgana catgnaccaa aaccaacttg gtc 283

<210> 2374  
 <211> 283  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2374

gannagtgc angcacgcgt acgtaagctc ggaattcggc tcgaggnaca gcacaaagcg 60  
 ggttactgtc tgttcaagct accatctctc tctctctttc ttagtgcctc cttgccagaa 120

gttaaaatgg cccaagaaac tttcctattc acatctgaat cagtgaacga ggggcaccct 180  
gacaagctct gtgaccagat ctccgatgct ntgctogatg catgcttgga gcaggaccct 240  
gacagcaagg ttgcctgtga aaactgcaac caagaccaan nat 283

<210> 2375  
<211> 302  
<212> nucleic acid  
<213> Glycine max  
<400> 2375

nanncgchang cacgcgtacg taagctcgga attcggctcg agctcgagcc gtggagtttg 60  
gagcgactga actaatcatt aatttgcact ncgctgtttc agcttcatca cccttctttt 120  
gcatcattta tatctcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180  
taaachnaggg tcaccccgac aagctgtncn accagatctc tgatgcagta cttgatgcgn 240  
gcctgancag gaccctgaca gcaagggttg ctgtgagaca tgcaccaagn cccaacatgg 300  
tc 302

<210> 2376  
<211> 270  
<212> nucleic acid  
<213> Glycine max  
<400> 2376

gcannngtnac gtaagctncg gaattcggct cgaggactta acaacagcac aaagcggggtt 60  
actgtgctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgaa 180  
caagctctgt gaccagatct ccgatgctgt gctcgatgca tgcttgagc aggaccctga 240  
cagcaagggtt gcctgtgana cctgcaccaa 270

<210> 2377  
<211> 312  
<212> nucleic acid  
<213> Glycine max  
<400> 2377

ngtcgcangc acgcgtacgt aagctcgga ttcggctcga gagaccaagc ccactcaac 60

cancacacca ntctactctg ctcttcttct acctttcaag tttctaaagt atnaagatgg 120  
cagagacatt cctattnacc tcagagtcag tgaacnaggg acacctgat caagctctgc 180  
gagccanac tccgatgctg tctctgacgg cttgccttga acaggaccca gacagcaagg 240  
ttgcctgcga aacatgcacc aagaccaact tggtcatggg cttcggagag atcaccncca 300  
aggccaacgt tg 312

<210> 2378  
<211> 328  
<212> nucleic acid  
<213> Glycine max

<400> 2378

agtcgcatgc acncgtacgt aagctcggaa tttcggctcg aggcattgtg gcaagtggac 60  
tagccagaag gtgcancncn gcaagtgtct tatgccattg gtgtgcnega ncctttgtnc 120  
tgtatttgtt gacacctatg gcaccgggaa gatccatgat aaggagattc tcaacattgt 180  
gaaggagaac ttgatttcan ncccggatg atctcccatc aaccttgatn tcaagagggg 240  
tgggaataac aggttcttga agatgctgca tatggacatt cggcagagag ncgnattcac 300  
aggggatggg cnangcccc ccaatggg 328

<210> 2379  
<211> 258  
<212> nucleic acid  
<213> Glycine max

<400> 2379

cagcacaang cnggttacng ncnngtcaag cnaccatctc tctctctctt tnttagtggc 60  
ctccttgcca gaagttaaaa tngcccaaga aactttcnta ttcacatctn aanccagtna 120  
acgaggggca cctgacaag ctctgtganc agatcnccga tgcngtgctc gntgcatnnt 180  
tggagcanga cctgacagc aaggttgcct gtgaaacctg naccaanacc aacatggnga 240  
ngggttcggg gagatcac 258

<210> 2380  
<211> 267  
<212> nucleic acid  
<213> Glycine max

<400> 2380

gtcgcanang angcgnacgt ncagctcgga attcggtctg aggtccggta tgatctccat 60

caaccttnga tctcaagagg ggtgggaata acagggttctt gaagactgct gcatatggac 120

acttcggcag agaggaccct gacttcacat gggaagtggg caagcccctc aagtgggaga 180

nggcctaagg ccattcattc cacngcaatg tgctgggagt ttttnagcgt tgcccttata 240

atgnctatta tccataactt tccacgt 267

<210> 2381

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2381

agtcnnatna acgcgtacgt aagctcggaa ttcggtctga gtacggctgc nagaagacga 60

cagaaggggg cagcgcttga tttgaggcca ggcaagcccc actcaaccac cacacctctc 120

ctcgttcacg ctaccctttt ctgctcttct tctacctttc aagtttttaa agtataaaga 180

tggcagagac attcctattt acctcagagt cgggtgaacga gggacaccct gacaagctct 240

gcgaccaaat ctccgatgct gtctctgacg cttgcctcga gcaggaccca gacagcaaag 300

ttgcctgcga a 311

<210> 2382

<211> 235

<212> nucleic acid

<213> Glycine max

<400> 2382

tttgaggcca ggcaagcccc actcaaccac cacacctctc ctcgttcacg ctaccctttt 60

actgctctnc ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt 120

tacctcagag tcggtgaacg agggacaccc tgacaagctc tgcgaccaa tctccgatgc 180

tgctctcgac gcttgctcgc agcaggaccc agacagcaaa gttgcctgcg aaaca 235

<210> 2383

<211> 168

<212> nucleic acid

<213> Glycine max

<400> 2383

ncngcncgng tacnggtacg cnagctcgga attcggctcg aggtgctggt gaccagggtc 60  
acatgttttg ctatgccact gatgaaaccc ctgaattgat gccattgagc catgttcttg 120  
caacaaaact cgggtgctcgt ctcaccgagg ttcgnaagaa cggtaacct 168

<210> 2384

<211> 156

<212> nucleic acid

<213> Glycine max

<400> 2384

tagtgcctcc ttgccagaag ttaaaatggc ccaagaaact ttctattca catctgaatc 60  
agtgaacgag gggcaccctg acaagctctg tgaccagatc tccgatgctg tgctcgatgc 120  
atgcttgag caggaccctg acagcaaggt tgctg 156

<210> 2385

<211> 278

<212> nucleic acid

<213> Glycine max

<400> 2385

ggccaggcaa gcccactca accaccacac ctctcctcgt tcacgctacn cttttctgct 60  
cttctctac ntttcaagtt ttaaaagtat aaagatggca gagacattnc ctatttacct 120  
canagtcggt gaacgaggga caccctgaca agctctgcga ccaaattctcc gatgctgtcc 180  
tcgacgcttg cctcgagcag gactcagaca gcaaagttgc ctgcgaaaca tggcaccaan 240  
accaattggt catggtcttc ggagaatcac gnccaagg 278

<210> 2386

<211> 278

<212> nucleic acid

<213> Glycine max

<400> 2386

gtcgcangca cgcgtacgtn agctcggnat tcggctcgag ggccaggcaa gcccactca 60  
accaccacac ctctcctcgt tcacgctacc cttttctgct cttctctac ctttcaagtt 120  
ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180



accctgacaa gctctgacgac caaatctccg atgctgtcct tcgacgcttg cctcgagcag 240  
gacccagaca gcaaagttgc ctgcgaaaca tgcaccaa 278

<210> 2387  
<211> 309  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2387

nngtcgcang cacgcgtacg taagctcgga attcggtcgc agtacggctg cnagaagacg 60  
acagaagggg gcagcgcttg atttgaggcc aggcaagccc cactcaacca ccacacctct 120  
cctgcgttca gctacccctt tctgctcttc ttctaccttt caagttttta aagtataaag 180  
atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc tgacaagctc 240  
tgcgacaaaa tctccgatgc tgtcctcgac gcttgccctg agcaggaccc agacagcaaa 300  
gttgccctgc 309

<210> 2388  
<211> 219  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2388

gctcggaatt cggctcgagc acaaagcggg ttactgtctg ttcaagctac catctctctc 60  
tctctttctt agtgccctct tgccagaagt taaaatggcc caagaaactt tcctattcac 120  
atctgaatca gtgaacgagg ggcaccctga caagctctgt gaccagatct ccgatgctgt 180  
gctcgatgca tgcttggagc aggaccctga cagcaaggt 219

<210> 2389  
<211> 314  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2389

gngtacgtna gctcggaatt cggctccgag ngatttgagg ccaggcaagc cccactcaac 60  
caccacacnt ctctcgttc acgctacccc tttctgctct tcttctacct ttcaagtttt 120  
aaaagtataa agatggcaga gacattccta ttacctcag agtcggtgaa cgagggacac 180

cctgacaagc tctgcgacca aatctccgat gctgtcctcg acgcttgctt cgagcaggac 240  
ccagacagcn aagttgcctg cgaaactgca ccaaaaccaa tnggtcaggt cttcggaat 300  
cngacaagcc acgt 314

<210> 2390  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
<400> 2390

ggaannntcg cangcacgcg tacgttagct cggaattcgg ctgagnact taacaacagc 60  
acaaagcggg ttactgtctg ttcaagctac catctctctc tctctttctt agtgctctct 120  
tgccagaagt taaaatggcc caagaaactt tcctattcac atctgaatca gtgaacgagg 180  
ggcacctgac aagctctgtg accagatctc cgatgctgtg ctgatgcat gttggagcag 240  
gacctgacag caaggttgcc tgtgaaacct gcaccaagac caanatg 287

<210> 2391  
<211> 281  
<212> nucleic acid  
<213> Glycine max  
<400> 2391

annantcgca ngcacgcgta cgtnagctcg gaattcngct cgagggccag gcnagcccca 60  
ctcaaccacc acacnactcc tenttencgc taccctttt ctgcnctct tctacctttc 120  
aagtttttaa agtatanaga tggcagagac attcctattt acctcagagt cggatgaacga 180  
gggacaccct gacaagctct gcgaccaaatt ctccgatgct gtgcttcgac gcttgctcgc 240  
agcaggacnc agacagcaaa gttgcctgcg aaacatgcac c 281

<210> 2392  
<211> 275  
<212> nucleic acid  
<213> Glycine max  
<400> 2392

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag caggcaagcc ccaactcaacc 60  
accacacctc tcctcggttca cgctaccctt ttctgctctt cttctacctt tcaagtttta 120

aaagtataaa gatggcagag acattcctat ttacctcaga gtcggtgaac gaggacaccc 180  
 tgacaagctc tgcgaccaa tctccgatgc tgtcctcgac gcttgcntcg agcaggaccc 240  
 agacagcaaa gttgcctgcg aaacatgcag caaaa 275

<210> 2393  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max

<400> 2393

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagt acggctgcga gaagaccaca 60  
 gaagggggca gcgcttgaga ccaagcccca ctcaaccacc acaccactct ctctgctctt 120  
 cttctacctt tcaagttttt aaagtattaa gatggcagag acattcctat ttacctcaga 180  
 gtcagtgaac gaggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga 240  
 cgcttgctt gaacaggacc cagacagcaa ggttgctgga aacatgnana agnccatntg 300  
 ggt 303

<210> 2394  
 <211> 189  
 <212> nucleic acid  
 <213> Glycine max

<400> 2394

cgtaagctcg gaattcggct cgaggtttca acaccttaat ttgcacacgc tgctttctca 60  
 gcttgagaaa tggcacaaga nacctttcta ttcacatctg aatctgtaaa cgagggtcac 120  
 cccgacaagc tgtgcgacca gatctctgat gcagtgcctg atgcgtgcct tgaacaggac 180  
 cctgacagc 189

<210> 2395  
 <211> 183  
 <212> nucleic acid  
 <213> Glycine max

<400> 2395

gctgcacgcg tacgtaagct cggaattcgg ctcgagctca agtttttgaa gtatagagat 60  
 ggcagagaca ttctatttta cctcagagtc agtgaacgag ggacaccctg acaagctctg 120

tgaccaaadc tctgatgctg tctctgacgc ttgcctcgaa caggacccag acagcaaggt 180  
tgc 183

<210> 2396  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
<400> 2396

gtcgcangca cgcgtacgtn nagctcggna ttccggctcgn cctctgagcc gaattcgggt 60  
cgagcaagcc ccaactcaacc accacaccnc tctctgttca cgcacncccc tttctgtctt 120  
tcttccacct ttcaagtttt aaaagtataa agatggcaga gacattcctt ttacctcaga 180  
gtcgggtgaac gagggacacc ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga 240  
cgcttgctc gagcaggacc cagacagcaa agttgctgc gaaacatgca cc 292

<210> 2397  
<211> 271  
<212> nucleic acid  
<213> Glycine max  
<400> 2397

gtcgcangca cgcgtacgtn agctcggnaa ttccggctcga gngatttgag gccaggcaag 60  
ccccactaca accaccacac ctctcctacg ttacgctac ccctttctgc tcttcttcta 120  
cccttcaagt tttaaaagta taaagatggc agagacattc ctatttacct cagagtcggt 180  
gaacgagggga caccctgaca agctctgcga ccaaattctc gatgctgtcc tcgacgcttg 240  
cctcgagcag gaccagaca gcaaagttgc c 271

<210> 2398  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
<400> 2398

cncgcgtacg taagctcgga attcgnctcg aggcagactt tcacaacagc acaaagcggg 60  
ttactgtctg ttcaagctac catctcgtc tcgctttctt agtgctcct tggccanaag 120  
tnaaaatggc ccaagaaact tncctattca catctgaatc agtgaacgag gggcaccctg 180

acaagctctg tgaccagatc tccgatgctg tgctcgatgc atgtcttgga gcaggaccct 240  
gacancaagg ttgcctgtga aactgcacca agaccaacat ggtgatg 287

<210> 2399  
<211> 307  
<212> nucleic acid  
<213> Glycine max  
<400> 2399

gcangcncgc gtncgtgagc tcggnttng gctcgnnggc caggcaagcn cactcancc 60  
accacacctc tntcgttca cgtacccct ttctgctctt cttctacctt tcangtttta 120  
anagtataaa gntggcagag acnttctnt ttncctcaga gtcggtgaac gagggacacc 180  
ctgacaagct ctgcgaccaa atctccgatg ctgtcctcga cgcttgctgc gagnanggac 240  
cnagacngca agttgcctgg gaaacatgca ccaggaccaa tttggtaatg gtctcgggaa 300  
aatcgng 307

<210> 2400  
<211> 291  
<212> nucleic acid  
<213> Glycine max  
<400> 2400

ngtcgctgca cgcgtacgta agctcggaat tcggctcgag gacttaacaa cagcaciaag 60  
cgggttactg tctgttcaag ctaccatctc tctctctctt tcttagtgcc tccttgccag 120  
aagttaaaat ggcccaagaa actttcctat tcacatctga atcagtgaac gaggggnacc 180  
ctganaagct cngngacnng nnntcngnng tngnncnng gngngctnga ggngggccct 240  
nacagcaagg ttgcctgtga aacctgcacc aagaccaaca tggatgatgt t 291

<210> 2401  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
<400> 2401

agtcgcangc acgcgtacgt aagctcgga ttggctcga gggccaggca agccccactc 60  
aaccaccaca cctctcctcg ttcacgtac cctttctgc tcttcttcta cctttcaagt 120

tttaaaagta taaagatggc agagacattc ctattttacct cagagtcggt gaacgagggg	180
caccctgaca agctctgcca cccaaatctc cgatgctgtc ctcgacgctt gncctcgagc	240
aggaccaga cagcaaagtt gcctgcgaaa catgcaccaa aaccaacttg gtcatggtct	300
tcgg	304

<210>	2402
<211>	302
<212>	nucleic acid
<213>	Glycine max
<400>	2402

anncangngt acgtnagctc ggaattcggg tcgagnacna cagnacaatg cgggttactg	60
tctgttcaag ctaccatcta ctctctcaact ttcttaacngc ctcccttgccc agaagttaaa	120
atgacccaag aatcttttct agtcacatct gaatcagcga acgaggngca ccttgacaag	180
ctccgtgacc agatctccga tgctgtgccc gatgcattct tggagcagga cccnacagca	240
agtttgctg ttaaacctgt nccaagacca acatggtgat ggtttgggaa anatcacaac	300
cn	302

<210>	2403
<211>	289
<212>	nucleic acid
<213>	Glycine max
<400>	2403

gtcgcangca cgcgtacgta agctcggaat tcggctcgag cacctctcct cgttcacgct	60
acccctttct gctctttctt tacctttcaa gttttaaaag tataaagatg gcagagacat	120
tcctattttac ctgagagtcg gtgaacngn gacaccctga caagctctgc gaccaaactc	180
ccgatgctgt cctcgacgct tgccctcgagc aggaccaga cagnaaagnt nccngcgaaa	240
canntacca aaaccaacnt ngnnanngnc atnggagaaa ncacgacca	289

<210>	2404
<211>	316
<212>	nucleic acid
<213>	Glycine max
<400>	2404

nnnanaacct tanaagtcgc angcacgcgt acgtaagctc ggaattcggc tcgaggctgc 60  
gagaagacga cagaaggggg cagcgcttga tttgaggccg ggcaagcccc actcaaccac 120  
cacacctctc ctcggttcacg ctaccccttt ctgctcttct tctacctttc aagttttaaa 180  
agtataaaga tggcagagac attcctattt acctcagagt cgggtgaacga gggnnaccct 240  
gacaagctct gcgaccaaatt ctccgatgct gtectcgacg cttgcctcga gcaggaccca 300  
gacagcaaag ttgcct 316

<210> 2405  
<211> 264  
<212> nucleic acid  
<213> Glycine max

<400> 2405

cgtcgcatgc acgcgtacgt aagctcggaa ttcggtcga gntttggagt ttggagcgac 60  
tgaactaatc attaatattgc actcgctgtt tcagcttcat gcacctttct tttgcatcat 120  
ttatatctct tgagaaatgg cacaagnac ctttctattc acatctgant ctgtaaacga 180  
gggtcacccc gacatgctgt ncgaccagat ctctgatgca gtacttgatg cgtgccttga 240  
acaggaccct gacagcaagg ttgn 264

<210> 2406  
<211> 308  
<212> nucleic acid  
<213> Glycine max

<400> 2406

nctagcacnc gtacgtaagc tcgggaattc anctcgann gcaagccccac tcaaccacca 60  
cacctctcct cggttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 120  
tataaagatg gcagagacat tcctatttac ctgagagtcg gtgaacgagg gacaccctga 180  
caagctctgc gaccaaattct gccgatgctg tcctgcgacg cttgcctcga gcaggaccca 240  
gacagcaaag ttgcctggcg aaacatgtac caaaaccaac ttggatcatg tnttcggaga 300  
aatcacga 308

<210> 2407  
<211> 331  
<212> nucleic acid

<213> Glycine max

<400> 2407

tgcttgacag cgtacgtaag ctcggaattc ggctcgaggt tgagaccaag acacactcgt 60  
tcatatatct ctncctgtct tctcttcttt ctacctctca agtttttgaa gtataaagat 120  
ggcagagaca ttcttattca cctcggagtc agtgaacgag ggacaccctg ataagctctg 180  
cgaccaaate tcgatgtctgt cctcgacgct tgccctgaac aggaccagat cagccangtt 240  
gcctgcgnaa acatgcacca agaccaattg gtcatggtct tcggagagat caccaccagg 300  
gccangntga cnncaagatc gtgcgtnaca c 331

<210> 2408

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2408

agtcgcaggg acgcgtacgt aagctcggna ntcggctcga ggcaagcccc actcaaccac 60  
cncancgntc ctcggttnac gctaccctt tctnggtctt ctntacctt tcangtttta 120  
aaagtntaaa gatggnagag acattcctat ttacctcaga gtcggtgaac gagggacacc 180  
ctgagaagct ctgcgaccaa atctccgatg tgctctcgac gcttgccctg agcaggaccc 240  
agacagcaaa gttgcctgcg aaacatgcac caaaaccaat tggnnaggtc ttcggagaat 300  
caggacaagg ccaaggtgan t 321

<210> 2409

<211> 242

<212> nucleic acid

<213> Glycine max

<400> 2409

ngcacgcgta cgtaagctcg gaattcggct cgaggcagac ttaacaacag caciaagcgg 60  
gntactgtct gttcaagcta ccactctctt ctctctttct tagtgccctc ttgccagaag 120  
ttaaaatggc ccaagaaact ttcttattca catctgaatc agtgaacgag gggcaccctg 180  
acaagctctg tgaccagatc tccgatgctg tgctcgatnc atgcttggag caaggccctg 240  
ac 242



<210> 2410  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2410  
  
 catgcacgcg tacgttagct cggaattcgg ctcgaggggt tactgtctgt tcaagctacc 60  
 atctctctct ctctttctta gtgcctcctt gccagaagta aaatggccca agaaactttc 120  
 ctattcacat ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc 180  
 gatgctgtgc tcgatgcagt cttggagcag gacctgaaag naaggttcct gtnaanttgc 240  
 acaancccaa tgggggggnt tttgggnagn nccacagng gggggggggn 289

<210> 2411  
 <211> 239  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2411  
  
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 tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
 aaaatggccc aaganacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180  
 aagctctgtg accagatctc cgatgctgtg ctcgatgcag gcttggagca ggaccctga 239

<210> 2412  
 <211> 249  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2412  
  
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 ctctgcttcc agcgagtgtt ctttcttcgt ttcaacacct taatttgcac acgctgcttc 120  
 ttcagcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg taaacgaggg 180  
 tcaccccgac aagctgtgcg accagatctc tgatgcagtg ctcgatgcgt gccttgaaca 240  
 ggaccctga 249

<210> 2413

<211> 121  
 <212> nucleic acid  
 <213> Glycine max

<400> 2413

tgtggcaagt ggactagcca gaaggtgcat tgtgcaagtg tcttatgcca ttggtgtgcc 60  
 cgagcctttg tctgtctttg ttgacaccta tggcaccggg aagatccatg ataaggagat 120  
 t 121

<210> 2414  
 <211> 273  
 <212> nucleic acid  
 <213> Glycine max

<400> 2414

tctnatgcac gogtacgtaa gctcggaatt cggctcgagg ccatttgga gttaggttct 60  
 gcacgctctg cttccagcga gtgttctttc ttcgtttcaa caccttaatt tgcacacgct 120  
 gcttcttcag cttgagaaat ggcacaagaa acctttctat tcacactgaa tctgtaaacg 180  
 agggtcaccc cgacaagctg tgcgaccaga tctctgatgc agtgcctgat gogtgccttg 240  
 aacaggaccc tgacagcaag gttgcctgtg aga 273

<210> 2415  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max

<400> 2415

gcangcacgc gtacgtaagc tcggaattcg gctcgaggca aaggagtgat ttggagtttg 60  
 gagcgactga actaatcatt aatttgcaact cgctgtttca gcttcatcac ccttgctttt 120  
 gcatcattta tatctcttga gaaatggcac aagaaacctt tctattcaca tctgaatctg 180  
 taaacnaggg tnaccccgac angcnntncg anccagatct ctgatgcagt acttgatgcg 240  
 tgccttgaac aggnccctga cngcaagggt cctggnnaga catgcaccag ggcnaacag 300  
 ggtaagggnc ttgg 314

<210> 2416  
 <211> 295  
 <212> nucleic acid

2416  
 2417  
 250  
 nucleic acid  
 Glycine max  
 2417  
 2418  
 206  
 nucleic acid  
 Glycine max  
 2418

<213> Glycine max

<400> 2416

cgtcgcangc acgcgtacgt nagctcggnn ttccggtcga nctcgagccg aatcggctcg 60  
 agtgaggcca ggcaagcccc actcaanac cacacttctc ctenttcacg ctaccccttt 120  
 actnctcttc ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt 180  
 tacctcagag tcggtgaacg agggacaccc tganaagctc tgcgaccana tctccgatgc 240  
 tgtcctcgac gttgcctcga gcaggaccca gacagcaaag ttgcctgcga aacat 295

<210> 2417

<211> 250

<212> nucleic acid

<213> Glycine max

<400> 2417

ggccaggcna gcccactca accaccacac nctnctctg ttcacgtac ccctttctgn 60  
 ctctttctct acctttcatg ttttaaaagt ataaagatgg cagagacatt cctatttacc 120  
 tcagagtcgg tgaacgaggg acanctgac aagctctgcg accaaatctc cgatgctgtc 180  
 ctcgacgctt gcncgagcag gaccagaca gcaaagttgc ctgngaaaca tgcnacaaaa 240  
 accaanttgg 250

<210> 2418

<211> 206

<212> nucleic acid

<213> Glycine max

<400> 2418

attaacttgg accttaagag ggggtggtcat aggttctca agacagctgc ttanggacac 60  
 tttggaaggg atgatgcaga cttcacctgg gaagttgtga agccactcaa gtcagagaag 120  
 cctcaagctt aagagtgttg ttaagttaat cactcccttc agtggatgtc ttgctgggtg 180  
 tggatgaata atttgcgtgt ttcatg 206

<210> 2419

<211> 152

<212> nucleic acid

<213> Glycine max

<400> 2419

nttcgcangc acgcgtacgt aagctcggaa ttengctcga ggaaactttc ctattcacat 60  
ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 120  
tcgatgcatg cttggannag gaccctgaca gt 152

<210> 2420

<211> 319

<212> nucleic acid

<213> Glycine max

<400> 2420

gtctcntgca cgcgtacgta agctcggaaat tcggntcgag gtttttgaag tatagagatg 60  
gcagagncat tcctatttac ctcagantcn gtgaacgagg nacnccctgn cangetgctg 120  
gtgaccaaat ctctgatgen gtcctcgacg cttgcctcga acaggacnca nacancangg 180  
ttgcctggng aancatgcac caaaaccaac ttgggtccatg gtcttcggag aaatcacgac 240  
caaggccnat gttgactacg agaagatagt gcgtgncacc tgcagagcnt cggctttgtn 300  
ctcaaacgat gtgggatgg 319

<210> 2421

<211> 262

<212> nucleic acid

<213> Glycine max

<400> 2421

gcagacttaa caacagcaca aagcnggta gtgtctgttc aagctaccat ctctctctct 60  
gctttcttag tgctccttg ccagaagtta aaatggcca agaaactttc ctattcacac 120  
ctgaatcagt gnacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 180  
tcgatgttgc ttggagcagg accctgacag caaggttgcn ngnaannnct gcaccangac 240  
aagnnnggtt ttgttgnacg ac 262

<210> 2422

<211> 231

<212> nucleic acid

<213> Glycine max

<400> 2422

gttgacgcg tacgtaagct cggaattcgg ctcgaggcag acttaacaac agcacaaagc 60  
gggttactgt ctgttcaagc taccatctct ctctctcttt cttagtgcct ccttgccaga 120  
agttaaaatg gcncaagaaa ctttcttatt cacatctgaa tcagtgaacg aggggcaccc 180  
tgacaagctc tgtgaccaga tctccgatgc tgtgctcgat gcattgcttg a 231

<210> 2423  
<211> 248  
<212> nucleic acid  
<213> Glycine max  
<400> 2423

ctaattgcacg cgtacgttag ctcggaattc ggctcgaggt gatttggagt ttggagcgac 60  
tgaactaatc attaatattgc actcgctgtt tcagcttcat cacccttctt ttgcatcatt 120  
tatatctctt gagaaatggc acaagaaacc tttctattca catctgaatc tgtaaacgag 180  
ggtcaccccg acangctgtn cnaccagatc tctgatgcag tacttgatgc gtgccttgan 240  
caggaccc 248

<210> 2424  
<211> 322  
<212> nucleic acid  
<213> Glycine max  
<400> 2424

tcgcaggcac gcgtacgtaa gctcggaatt cggctcgagn cagacttaac aacagcacia 60  
agcgggttac tgtctgttca agctaccatc tctctctctc ctttcttagt gcctccttgc 120  
cagaagttaa aatggcccaa gaaactttcc nattcacatc tgaatcagtg aacgaggggc 180  
accctgaaca agctctgtga ccagatnctc cgatggctgt gctcgatgnc atgcttggag 240  
caggaccctg acagcaaggt tgctgtgaa acctgcacca ggaccaacat ggtgatgggt 300  
ttcgagagat cacaaccaag gc 322

<210> 2425  
<211> 317  
<212> nucleic acid  
<213> Glycine max  
<400> 2425

tcgcangcac ncgtacgtna gctcgggaatt cggctcgagn ngccaggcaa gcccactca 60  
 accaccacac ctctcctcgt tcacgctacc cttttctgct cttctttctac ctttcaagtt 120  
 ttaaaagtat aaagatggca gagacattcc tatttacctc agagtcggtg aacgagggac 180  
 accctgacaa gctctgcgac caaatctccg atgctgtcct cgacgcttgc ctcgagcagg 240  
 acccagaaca gcaaagttgc ctgcgaaact ggcaccaaca ccaattggtc atggtontcg 300  
 gagaaatcnc gaccagg 317

<210> 2426  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max

<400> 2426

angcacgcgt acgtaagctc ggaattcggc tcgagcggct gcgagaagac gacagaaggg 60  
 ggcagcgctt gatttgaggc caggcaagcc ccacttcaac caccacacct ctctcgttc 120  
 acgtaccccc tttctgctct tcttctacct ttcaagtttt aaaagtataa agatggcaga 180  
 gacattccta ttacctcag agtcggtgaa cgagggacac cctgacaagc tctgcgacca 240  
 aatctccgat gctgtcctcg acgcttgctt cgagcaggac ccagaca 287

<210> 2427  
 <211> 347  
 <212> nucleic acid  
 <213> Glycine max

<400> 2427

cncnnnang tcgcatgcac gcgtacgtaa gctcgggaatt cggctcgaga cggctgcna 60  
 gaagacgaca gaagggggca gcgcttgatt tgaggccagg caagccccac tcaaccacca 120  
 cacctctcct cgttcacgct acccctttct gctcttcttc tacctttcaa gttttaaaag 180  
 tataaagatg gcagagacat tcctatttac ctcagagtcg gtgaacgagg gacacctga 240  
 caagctctgc gaccaaactt ccgatgctgt cctcgacgct tgccctcgagc aggaccaga 300  
 cagcaaattg cctgcgaaac atgcaacaaa aacaanttgt canggnc 347

<210> 2428  
 <211> 288  
 <212> nucleic acid

<213> Glycine max

<400> 2428

gntcangcac gcgtacgtaa gctcggaatt cngctcgagc annagcataa agcgggttac 60  
tgtctgttca agctacncat ctctctctct cttncctagt gcctccttnc cagaagttan 120  
natggcccaa gaaactttcc tattcacatc tgaatcagtg aancgagggg caccctgaca 180  
agctctgtga ccagatctcc gatgctgtgc tcgatgcatg cttggagcag gangngacag 240  
canggttgcc tgtgaaacct gcaccaagan caacatggtg atgntttt 288

<210> 2429

<211> 226

<212> nucleic acid

<213> Glycine max

<400> 2429

ctgcacnctg acgtacgctc ggcaacnctg tccactnaac caccacacct cncctcgtt 60  
cacgctaccc ctttctgctc ttncttctac ctttccaagt tttaaaagtn taaagatggc 120  
agagacattc ctatttacct cagagtcggt gaacgagggg caccctgaca agctctgcga 180  
ccaaatctcc gatgctgtcc tcgacgcttg cctcgagcag gaccca 226

<210> 2430

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2430

gcacgcgtac gtaagctcgg aattcggtc gagtacggt cgagaagacg acagaagggg 60  
gcagcgcttg atttgaggcc aggcaagccc cactcacacc accacacctc tctcgttca 120  
cgctacccct ttctgctctt cttctacctt tcaagtttta aaagtataaa gatggcagag 180  
acattcctat ttacctcaga gtcggtgaac gagggacacc ctgacaagct ctgcgaccaa 240  
atctccgatg ctgtcctcga cgcttgcttc gagcaggacc cagacag 287

<210> 2431

<211> 164

<212> nucleic acid

<213> Glycine max

<400> 2431

gcgtangtaa gctcggaatt cggctcgagg tcatatgttc ggctatgna ctgacgagac 60  
 tcccagagctc atgcccttga gccatgtcct tgccacgaag ctcggtgna agctcancga 120  
 ggttcggaan aacggganat gcccttggnt gannectnnt ggca 164

<210> 2432

<211> 292

<212> nucleic acid

<213> Glycine max

<400> 2432

gtcgcangca cgcgtacgta agctcggaat tcggctcgag nccatttggg agttagggtc 60  
 tgcacgctct gcttcacgag agtggtcttt ctctgtttca acaccttaat ttgcacacgc 120  
 tgcttcttca gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa 180  
 cgaggggtcac cccgacaagc nnncgaccag atngcnnang cagtgcgcga ngngngnct 240  
 nnacaggacc cngncagcaa ggcnngctgn nagacangca ncaagaccaa ca 292

<210> 2433

<211> 97

<212> nucleic acid

<213> Glycine max

<400> 2433

ccaagaccaa catggtcatg gtctctggag anctcacaac caaggccacc gtagactang 60  
 agaagattgt ccgtgacaca tgccgcgaaa ntggata 97

<210> 2434

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 2434

catgcacgcg tacgtaagct cggaattcgg ctcgagtacg gctgcgagaa gacgacagaa 60  
 gggggcaccg cttgagcaga cttacaaca gcacaaagcg ggttactgtc tgttcaagct 120  
 accatctctc tctctctttc ntagtgtccc ttgccagaag ttaaaaatgg cccaagaaac 180  
 tttcttattc acatctgaat cagtgaacga ggggcacctg acaagctctg tgaccagatc 240





tagtcatggt ggtgtttttg gctgtgaatt tgctcatatg tgctaattat gtgtttcttgt 180  
 ttgatgttac tctacagaag ttaaaatggc ccaagaaact ttcctattca catctgaatc 240  
 agtgaacgag gggcaccctg aacaagctct gtgaccagat ctccgatgct gtgctcgatg 300  
 catgcttgga gcagg 315

<210> 2438  
 <211> 121  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2438

attaacttgg accttaagag ggggtggcat aggttcctca agacagctgc tnatggacac 60  
 ttggaaggg atgatgcaga cttcacctgg gaagttgtga agccactcaa gtcagagang 120  
 c 121

<210> 2439  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2439

gtcgcacgca cgcgtacgta agctcggaat tcggctcgag caggcaagcc ccaactcaacc 60  
 accacacctc tctcgttca cgtaccctt ttctgtctt cttctacctt tcaagtttta 120  
 aaagtataaa gatggcagag anattcctat ttacctcaga gtcggtgaac gagggacacc 180  
 ctgacaagct ctgcancca aatctccgat gctgtcctcg acgcttgctt cgagcaggnc 240  
 ccagacagcc aaagttgcct gcgaaacang cagcnaaacc aacttggtc 289

<210> 2440  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2440

cntcangnac gcgtacgtaa gctcggaatt cgggctcgag nttgaggcca ggcaagcccc 60  
 actcaaccac cacacctctc ctcgttcacg ctaccctttt ctgctcttct tctacctttc 120  
 aagtttttaa agtataaaga tggcagagac attcctattt acctcagagt cgggtgaacga 180

gggacaccct gacaagctct ggggaccaa tctccgatgc tgcctcgcac gcttgccntn 240  
gagcaggacc cnganagcaa antngcttgg gaaanttgcn caaaaacccat ttgggtnnng 300  
gtntgngnaa 310

<210> 2441  
<211> 283  
<212> nucleic acid  
<213> Glycine max

<400> 2441

angcacngt acgtnagctc ggaattcggc tcgagcttaa caacagcaca aagcgggtta 60  
ctgtctgttc aagctaccat atctctctct ctnttcttag tgcctncott gccanaagtt 120  
aaaatggnc ntgaaacttt ccnattcaca tntnnatcag tgaacgaggg gcaccccgac 180  
aagctctntn atagatcngg gtngncagtg ctagatgnat gnttggagca ngancctnan 240  
agcnaggntn cctgtgaaac cnggcacna gaccaacatg gtn 283

<210> 2442  
<211> 240  
<212> nucleic acid  
<213> Glycine max

<400> 2442

tgaggccagg caagccccac tcaaccacca canatnantic ctggttncac gctacccctt 60  
tctnctctt cttctacctt tcangtttta anagtataaa gatggcagag acattcctat 120  
ttacctcaga gtcggtgaag agggacaccc tgacaagctc tgcgaccaa tctccgatgc 180  
tgtcctcgac gcttgctcgc agcaggaccc agacagcaaa gttgntggaa acatgcacca 240

<210> 2443  
<211> 296  
<212> nucleic acid  
<213> Glycine max

<400> 2443

tcgatgcac gcgtacgtaa gctcggaatt cggctcgagg gtttttgaag actgctgcct 60  
atggacactt tggaagagaa gaccctgact tcacatggga agtgggtcaaa cccctcaagt 120  
gggagaaggc ctaagtaant cattccactg ctctatgctg gaagtttttt gagcgttgcc 180

cttataatat gtctaataatc cataactttc cacgtctctt acnctgtgtg tttctctcct 240

cttctctcta ttttggttatt tgtatgttct tttgtaattt ttacgtgatc aactaa 296

<210> 2444

<211> 287

<212> nucleic acid

<213> Glycine max

<400> 2444

gtcgcangca cgcgtacgta agctcggaat tccgctcgag ngtgccata ggttctctca 60

gacagctgct tatggacact ttggaaggga tgaccctgac ttcacctggg aagttgtgaa 120

gccactcaag tctgagaagc ctcaagctta agattgttgt gaagttaatc actcccttca 180

atggatgtct tgctaggtgt ggatgaataa tttgcgtgtt ccatgactac tactacttca 240

ttcataggtc taatgtcatc tcatcaatac ttaaactgtt ttttttt 287

<210> 2445

<211> 185

<212> nucleic acid

<213> Glycine max

<400> 2445

agcagactta acaacagcac aaagcgggtt actgtctgtt caagctacca nnnnnnnnnn 60

nnnnnnntag tgctctcttg ccagaagtta aaatggccca agaaactttc ctattcacat 120

ctgaatcagt gaacgagggg caccctgaca agctctgtga ccagatctcc gatgctgtgc 180

togat 185

<210> 2446

<211> 227

<212> nucleic acid

<213> Glycine max

<400> 2446

ngtcatatgca cgcgtacgta agctcggaat tccgctcgag gttnggttct gcacgctctg 60

cttcacagca gtgttctttc ttcgtttcaa caccttanat ttgcacacgc tgcttcttca 120

gcttgagaaa tggcacaaga aacctttcta ttcacatctg aatctgtaaa cgaggggtcac 180

cccagacaagc tgtgcgacca gatctctgat gcagtgtctg atgcgtg 227

<210> 2447  
 <211> 98  
 <212> nucleic acid  
 <213> Glycine max

<400> 2447

ccttttcagg gaaggaccct accaagggtg acagaagtgg tgcctatatt gtaaggcagg 60  
 ctgcaaagag tgtcgtggca aatggccttg ctagaagg 98

<210> 2448  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max

<400> 2448

gtcgcngcac gcgtacgtna gctcggaatt nggctcgagc tcgagccgca acagcacaaa 60  
 gggggttact gtctgttcaa gctaccatct ctctctcact ttcttngtgc ctcttgcca 120  
 gangttanna tggcccaaga aactttccta ttcacatctg aatcagtga cgagggggcac 180  
 ncnnnacaag ctctgtgacc agatctccga tgctgtgcta cgatgcatgc ttggagcang 240  
 naccctgaca gcnaagttgc ctgtgaaacc tggcaccaag ancaacatgg tgatggtttt 300  
 cgga 304

<210> 2449  
 <211> 266  
 <212> nucleic acid  
 <213> Glycine max

<400> 2449

gtcgcgatgca cgcgtacgta agctcggaat tcggctcgag gtgatttga gtttgagcg 60  
 actgaactaa tcnttanttt gactcgtg tttcagcttc atcacccttc ttttgcata 120  
 tttatatctc ttgagaaatg gcacaagaaa cttttctatt cacatctgaa tctgtaaacg 180  
 agggtcaccc cgacangctg tgcgaccaga tctctgatgc agtacttgnn gcgngcctna 240  
 aaggncacca ncancaaggt cgcctg 266

<210> 2450  
 <211> 159  
 <212> nucleic acid

<213> Glycine max

<400> 2450

toggaattcg gctcgagaac agcacaaagc gggttactgt ctgttcaagc taccatctct 60  
ctctctcttt cttagtgcct ccttgccaga agttaaaatg gccaagaaa ctttcctatt 120  
cacatctgaa tcagtgaacg aggggcaccc tgacaagct 159

<210> 2451

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2451

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ccanaccact ctctctgtct ttcttctacc tttccaagtt tntaaagtat taagntggca 120  
gagacantcc nanntanctc agagncngng nangngggnc ancctgnan gcgctncgac 180  
naatctneca tgtgtctctg acgcttgctt tgaacaggac ccagacagca aggttgcttg 240  
cgaaacatgc accaaganca attggtcatg gtcttcggag agatcacca 289

<210> 2452

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 2452

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acgacgagac tgtcaccaat gangaaattg ntgntgacct tcaaanagca tgtgatcaan 120  
cctgngatnc cngngaantn nctnatnagn agnncanttt ccnattngaa cnttaaggc 180  
ggtttggaac tggttggccn nnaggggcna ngctgggtct ccggggncga aaagancctt 240  
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<210> 2453

<211> 181

<212> nucleic acid

<213> Glycine max

<400> 2453

gnngctgcac gcgtacgtna gctcgggaatt cggctcgagt aacaacagca caaagcgggt 60  
 tactgtctgt tcaagctacc atctctctct ctctttctta gtgcctcctt gccagaagtt 120  
 aaaatggccc aagaaacttt cctattcaca tctgaatcag tgaacgaggg gcaccctgac 180  
 a 181

<210> 2454  
 <211> 268  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2454

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 gctctgcttc cagcgagtgt tctttcttcg tttcaacacc ttaatttgca cacgctgctt 120  
 cttcagcttg aggaatggca caagaaacct ttctnttcac atctgantct gtanacgang 180  
 gtcacccoga caagntgtgc gaaccagatn ctctgatggc agtgctcatg cgtgcgctga 240  
 ncaggacct gacagcaagg ttgcctgn 268

<210> 2455  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2455

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 cgncanangg aaaaaacgaa ngcgcnaccc cattcngctc ntcttcnacc tnnrangann 120  
 naagagnnta aagatggcag agacattcct atttacctca gagtacggtg aacgagggac 180  
 accctgacaa gctctgcgan ccaagtctcc gatgctgtcc tcgncgcttg cctcgagcag 240  
 gaccagaca gcaaagttgc gcagcganac atgcancaag ncgnattggn ccatgggtt 298

<210> 2456  
 <211> 154  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2456

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gaggggaaga cccaagtacg cgttgagtat tacaatgaca atgggtgccag ggttcctatt 120  
ccgtgtacac accgtgctaa tttccacaca acat 154

<210> 2457  
<211> 284  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2457

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tggtcaagct accatctctc tctctctttc ttagtgccctc cttgccagaa gttaaaatgg 120  
cccaagaaac tttctatttc acatctgaat cagtgaacga ggggcaccct gacaagctca 180  
gagaccagaa nancgangcn gngcacgacg caagtccttg agcaggaccc agtacagcaa 240  
ggnnnnncagt gnaaccngta ccaagaccaa caggtgatgg tcct 284

<210> 2458  
<211> 213  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2458

gtencangca cacgtacgta agctcggaat tcnnctcgag ncaagcccca ctcaaccacc 60  
acacctctcc tegtacacgc tacccttttc tgctcttctt ctanctttca agtttttaaaa 120  
gtataaagat ggcagagaca ttcctattta cctcagagtc ggtgaacgag ggacaccctg 180  
acaagctctg cgaccaacan ctccgatgct gtc 213

<210> 2459  
<211> 217  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2459

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ctgtctgttc aagctacctt ctntctctct ctttcttagt gcctccttgc cagaagttaa 120  
aatggnccaa gaaactttcc tattcacatc tgnatcagtg ancgangggc accctgacaa 180  
gctctgtgac cagatctccg atgctgtgct cgntgca 217



<210> 2460  
 <211> 233  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2460  
  
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 tggagcgact gaactaatca ttaatttgca cttcgctgtt tcagcttcat cacccttctt 120  
 ttgcatcatt tatactctctt gagaaatggc acaagaaacc tttctattca catctgaatc 180  
 tgtaaacgag ggtcaccocg acangctgtg cgaccagatc tctgangnag nat 233

<210> 2461  
 <211> 202  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2461  
  
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 ccacacctct cctcgttcac gctaccctt tctgctcttc ttctaccttt caagttttta 120  
 aagtataaag atggcagaga cattcctatt tacctcagag tcggtgaacg agggacaccc 180  
 tgacaagctc tgcgaccaa tt 202

<210> 2462  
 <211> 196  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2462  
  
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 cagcacaaaag cgggttactg tctgttcaag ctaccatnct ctctctctct ttacttagtg 120  
 cctccttgcc agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga 180  
 acgaggggca ccctga 196

<210> 2463  
 <211> 323  
 <212> nucleic acid  
 <213> Glycine max

<400> 2463

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gttactgtct gttcaagcta ccattctctct ctctctttct tagtgcctcc ttgccagggtg 120  
ctgccactct ctctttctct ctcttcatcc ttctgttggg ttggttgtgg agtgttgttt 180  
tctgttgtgc acgtgttgtc attttttacc ctgccacag atctgaagtg ttcaagtttg 240  
gattttgtgc ttctggaagt taaaatggcc caagaaactt tcctattcac atctgaatca 300  
gtgaacgagg ggcaccctga caa 323

<210> 2464

<211> 132

<212> nucleic acid

<213> Glycine max

<400> 2464

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gttgagtaac tacaatgaca atggtgccat gggtccagtt cgtgtccaca ctgtcctaata 120  
ttcccacaca ac 132

<210> 2465

<211> 189

<212> nucleic acid

<213> Glycine max

<400> 2465

gggagttagg ttctgcacgc nctgnngcca gcgagtgtgc tntcttcgtg tcaacacctg 60  
aatttgcann acgctgcgnc tgcagcttga gaaatggcac aagaaaccnn gctatncana 120  
tctgaatcgg taaacgaggg tcacnncgac aagctgtggg accagatctc tgatgcagtg 180  
ctcgatgcg 189

<210> 2466

<211> 138

<212> nucleic acid

<213> Glycine max

<400> 2466

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catatgttcg gctatgccna ctgacgagnn ctcccagagct catgcccttg agccatgtnc 120  
 cttgccacga agcttcgg 138

<210> 2467  
 <211> 341  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2467

ncctctnanc ntttacatcg anacnacgta cgtnacgtcg gnattcggct cgagcagact 60  
 taacaacagc acaaagcggg ttactgtctg ttcaagctac catctctctc tctctttctt 120  
 agtgccctcc ttgccaggtg ctgccactct ctctttctct ctcttcatcc ttctgttggg 180  
 ttggttgtgg agtggtgttt tctgttgtgc acgtgttgtc attttttacc ctgccacag 240  
 atctgaagtg ttcaagtttg gattttgtgc ttctggaagt taaaatggcc caagaaactt 300  
 tcctattcac atctgaatca gtgaacgagg ggcaccctga c 341

<210> 2468  
 <211> 273  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2468

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 tctccctata aatggcaact caatgcccc cttagaactc gcagcgcttg atttgaggcc 120  
 aggcaagccc cactcaacca ccacacctct cctcgttcac gctacccctt tctgctcttc 180  
 ttctaccttt caagttttta aagtataaag atggcagaga cattcctatt tacctcagag 240  
 tcggtgaacg agggacaccc tgacaagctc tgc 273

<210> 2469  
 <211> 181  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2469

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 tcgttcacgc tacccttttc tgctcttctt ctacctttca agttttaaaa gtataaagat 120

ggcagagaca ttctatttta cctcagagtc ggtgaacgag ggacaccctg acaagctctg 180

c 181

<210> 2470

<211> 305

<212> nucleic acid

<213> Glycine max

<400> 2470

gtcgcattgca cgcgtacgta agctcggaat tcggctcgan gnagacttaa cancagnnca 60

aagcgggtta ctgtctgntc angctaccat ctctcnctct ntttcttagt gcctccntgc 120

cagnagttnn aatggcccaa gnnacttten tantcacatc tgantcnntg aacgaggggc 180

accngataa gctctgtgan cagatctccg atgctgtgct ccgatgnatg cttggagcng 240

gnnnctgnca gcnaggntgn ctgtgnaach tgcacnangn ncancatggt gatggntttc 300

ggnga 305

<210> 2471

<211> 199

<212> nucleic acid

<213> Glycine max

<400> 2471

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agcgggttac tgtctgttca agctaccatc tctctctctc tttcttagtg cctccttgcc 120

agaagttaaa atggcccaag aaactttcct attcacatct gaatcagtga acgaggggca 180

ccctgacaag ctctgtgat 199

<210> 2472

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2472

cgcattgcacg cgtacgtaag ctcggaattc ggctcgagcc ttgatctcaa gaggggtgga 60

aatggcaggt tcttgaagac tgctgcatat ggacactttg gcagagatga ccctgacttc 120

acatgggaag tggatgaagcc actcaagggg gagaaggtag ctgcttaact aaaaggggtt 180

ccaacactct tggcaaggga cttttgcact actactggct tcttattatc tgattgctaa 240  
aattttctct atgtttcctt ccctctact caattctggt ttttttttnc ngatattttt 300  
tatgaatttc cccctttttt ttgtgta 327

<210> 2473  
<211> 256  
<212> nucleic acid  
<213> Glycine max  
<400> 2473

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ctgacttcac atgggaagtg gtgaagccac tcaaggggga gaaggtagct gcttaactaa 120  
aaggggttcc aacactcttg gcaagggact tttgcactac tactggcttc ttattatctg 180  
attgcnaaaa tttnctctat gnntccttcc ctcttactca attctgtttt tntttttctg 240  
tatttttnat gaattc 256

<210> 2474  
<211> 214  
<212> nucleic acid  
<213> Glycine max  
<400> 2474

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catgggaagt ggtgaagcca ctcaaggggg agaaggtagc tgcttaacta aaaggggttc 120  
caacactctt ggcaagggaac ttttgcacta ctactggctt cttattatct gattgctaaa 180  
attttctcta tgtttccttc cctcttactc aatc 214

<210> 2475  
<211> 206  
<212> nucleic acid  
<213> Glycine max  
<400> 2475

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catgggaagt ggtgaagcca ctcaaggggg agaaggtagc tgcttaacta aaaggggttc 120  
caacactctt ggcaagggaac ttttgcacta ctactggctt ctattatctg attgctaaaa 180

ttttctctat gtttccttcc ctctta

206

<210> 2476  
<211> 311  
<212> nucleic acid  
<213> Glycine max

<400> 2476

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tggacacttt ggcagagatg accctgactt cacatgggaa gtggtgaagc cacttcaagg 120  
gggagaagggt acctgcttaa ctaaaagggg ttccaacact cttggcaagg gacttttgca 180  
ctactactgg cttcttatta tctgattgct aaaattttct ctatgtttcc tccctcttac 240  
tcaattctgt ttttttttnt ctgtnttttc tnatgaattt cccctttttt tttgggnact 300  
ngnatgtgtt c 311

<210> 2477  
<211> 300  
<212> nucleic acid  
<213> Glycine max

<400> 2477

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natgacctg acttcacatg ggaagtgggtg aagccacttc aagggggaga aggtacctgc 120  
ttaactaaaa ggggttccaa cactcttggc aagggaactt tgactacta ctggcttctt 180  
attatctgat tgctanaatt ttctctatgt ttcttccct ctactcaat tctntttttc 240  
nttttctgta tttntttatg aatttcccc tttntntgn gnacttgtn gngtnctnnc 300

<210> 2478  
<211> 291  
<212> nucleic acid  
<213> Glycine max

<400> 2478

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ncaaacgagg gccaccccgga caagatctgt gaccaggttt ctgatgccat cctcgatgct 180

tgcttggagc aagaccaga gagcaaggtt gcctgcgaga cctgtacaaa aaccaacatg 240  
gtcatgggtct ttggggagat cacaaccaag gccaaaggtga actacgagaa t 291

<210> 2479  
<211> 308  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2479

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gaaggggttc ctcttcaatt tcgcatcttc tccttctcat tccaacttcc aaaatacaca 120  
cacacgatgg aaaccttctt cttcacctca gaatctgtaa acgagggcca ccccgacaag 180  
atctgtgacc aggtttctga tgccatcttc gatgcttgct tggagcaaga cccagagagc 240  
aaggttgctt gcgagacctg tacaaaaacc aacatgggtca tggctcttgg ggagatcaca 300  
accaaggc 308

<210> 2480  
<211> 262  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2480

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attgtgttga cgcnggactg aatgaactaa tggagtctan aggtgggaaa nagaagtcnn 120  
nnnnnnnnnn nnnnnaatca ttgttctacg aagctccctt cggatacagc atnngaagac 180  
gttagacca aaggtggaat caagaaattc agatctgctg cttactccaa cgtatatntt 240  
cttctgatgc agtgattctg ta 262

<210> 2481  
<211> 420  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2481

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gtaagntcgg aattcggctc gagctcgagc cgcaaaaatt cgggaagact agtgactgtg 120

gttcacctag atcaactcta aagtgtctga atgaggaaga tgaggaagag tagtttcctt 180  
aagtgtcttt attattctgt ccttgtgaaa ataagtctgg ttttccagat acgttattgt 240  
ttttctttgt tgtctttttt agcttctgtt agagaccatt tgggcattta gacctttatt 300  
gtttctatta ccatttgaac atcgaatgga ttaataaatc actttgtttg cgtgcaaaaa 360  
aaannacana tctttcnana aanaaaaaaa annaaaaana acanaaaaaan aaaaaaaaaan 420

<210> 2482  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
<400> 2482

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cattgtccaa agaccaattg gacgagattc ttaaaccagc agagtgcact attgtngcat 120  
cactttcaaa tgattatgtt gactcttatg ttctgtcaga gtcaagcctg ttctgtctatc 180  
cttataaaaat tatcatcaaa acttgtggga ctaccaaatt gcttctgtcc atccctgtcc 240  
attctcaagt tgggctgatg ctcttgacat agctgtgaaa tctgtga 287

<210> 2483  
<211> 288  
<212> nucleic acid  
<213> Glycine max  
<400> 2483

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ccaaagacca attggacgag attcttaaac cagcagagtg cactattgtt gcatcacttt 180  
caaatgatta tgttgactct tatgttctgt cagagtcaag cctgttcgtc tctcctataa 240  
aattatcatc aaaacttgtg ggactaccaa attgcttctg tccatccc 288

<210> 2484  
<211> 306  
<212> nucleic acid  
<213> Glycine max  
<400> 2484



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 caaagg 306

<210> 2485  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max

<400> 2485

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 aatttgaccc ttgcggatat tcaatgaatg gaatagaagg gagtgctata tccaccatcc 120  
 atgtcactcc tgaagatggg ttcagttacg caagttttga agctgttggt tatgacttta 180  
 atgacatggc tctaggtgaa cttgtggaaa ggatttttagc tgcttttgtc cagcagagtt 240  
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 gatactactg tggg 314

<210> 2486  
 <211> 476  
 <212> nucleic acid  
 <213> Glycine max

<400> 2486

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 ctgcgatagt ttctctactg ttacatggcc atggcgggtt ccgcaattgg ttttgaaggt 180  
 ttcgagaaaa ggttggaaat atcctttttc cagccgggac tttttgctga ccctgaagga 240  
 aggggtctaa gggctcttac aaaatcccag ttgggtgaga ttctaacc accagctgttc 300  
 accattgttt cttegtctaa aaacgataat gtcgactcct atgttctatc tgagtccagc 360  
 ctctttgttt atgcctacaa gatcatcatc aaaacctgtg gtactactaa gctattgctt 420

gcaatcccac ccatattgaa gttcgtgaa atgctttccc ntaatgttaa gtcngt 476

<210> 2487  
 <211> 510  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2487

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 cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag ctgccatgat 180  
 gactgttaat tccggcatta gaaaaattct tccagattcc gagatttgtg actttgactt 240  
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 ggtgaatctg aacgaaatgg gttcaagang gtattggcat gtttcctncc aactgagttt 420  
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 taagggatac tgnctgaaaa gaggaacccc 510

<210> 2488  
 <211> 560  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2488

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 gaaaaattct tccagantct gagatttgtg actttgactt tgagccatgt ggttattcaa 180  
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 ttaagagggg attggcatgt tttctcccaa ctgagttctc tgttgcagtt catgtggatg 360  
 gtgcaagcaa gttgtttgat cagacgtgtn ttctggatgt taagggatac tgtcgcaag 420  
 agaggagccc acgaaagggg ttgggaatgg gtggnnntct tggctaccaa aaaaantgcc 480  
 aaagacttgg gaactggggg tcaactagan ccaactctga aangntggaa aagaaggaag 540

atgaagaaag agtagttttt

560

<210> 2489  
<211> 485  
<212> nucleic acid  
<213> Glycine max  
<400> 2489

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ctgctgtttc taccattcat gttaccccag aagatgggtt cagttatgca agctttgaga 180  
ctgttgggta tgacttcaaa gtggtgaatc tgaacgaaat ggtaagagg gtattggcat 240  
gttttctccc aactgagttc tctgttgagc ttcattgtga tgggtgcaagc aagttgtttg 300  
atcagacgtg ttttctggga tgttaaggga tactgtcgcg aagagaggag ccacgaaggg 360  
cttggaatgg gtggttctct tgtctaccaa aaatttgcca agacttgtga ctgtggttca 420  
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aagtg 485

<210> 2490  
<211> 339  
<212> nucleic acid  
<213> Glycine max  
<400> 2490

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ctgccatgat gactgttaat tccggcatta gaaaaattct tccagattcc gagatttgtg 120  
actttgactt tgaaccatgt gggtattcaa tgaactctgt tgaaggcgct gctgtttcta 180  
cgattcatgt taccocagaa gatggtttca gttatgcaag cttcgaaact gttggttatg 240  
acttcaaagc ggtgaatctg aacgaaatgg ttcagagggt attggcatgt ttcctcccaa 300  
ctgagttttc tgttgcagtt catgtggatg gtgcaagca 339

<210> 2491  
<211> 412  
<212> nucleic acid  
<213> Glycine max

<400> 2491

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aatgggttaa gaggggtattg gcatgttttc tcccaactga gttctctgtt gcagttcatg 120  
tggatgggtgc aagcaagttg tttgatcaga cgtgttttct ggatgttaag ggatactgtc 180  
gcgaagagag gagccacgaa gggcttggaa tgggtgggtc tcttgtctac caaaaatttg 240  
ccaagaacttg tgactgtggt tcacctagat caactctgaa gtgctggaaa gaggaagatg 300  
aagaagagta gttttcttaa gtgtctttat tatgtccttg cgaaaataag tccggttttc 360  
cagacagtga ttgtttntct ttggtgnttt ttnccttnta tgtagacca tg 412

<210> 2492

<211> 504

<212> nucleic acid

<213> Glycine max

<400> 2492

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cgtacgtaag ctcggaattc ggctcgagtg catgcaacca tagttttatt aggatttttt 120  
cttctttgtt ttcaattagg tttttttgtt gctctccttc aaactccatc tttccaaatc 180  
ctctctttgc gattgtgttt tgatctgctt cctactgcga tagtttctct actgttacat 240  
ggccatggcg gtttccgcaa ttgggttttg aggtttcgag aaaagggttg aaatatcctt 300  
tttccagccg ggactttttg ctgaccctga aggaaggggt ctaagggtc ttacaaaatc 360  
ccagttgggt gagattctaa caccagctgc ttgcaccatt gtttcttcgc tcaaaaacga 420  
taatgtcgac tctatgttc tatctgagtc cagcctcttt ggttatgcct acaagatcat 480  
catcaaaacc tnggggacta ctaa 504

<210> 2493

<211> 347

<212> nucleic acid

<213> Glycine max

<400> 2493

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ttttgtgac cctgagggaa tgggtttaag agctcttgca nagtcccagt tggatgagat 120

acttacaccg gctgcttgca ccattgtttc atctctcaga aatgatcatg tcgactccta 180  
 tgttctgtct gagtccagtc tctttgttta tgcctacaag atcatcatca aaacctgtgg 240  
 tactacaaag ctactgcttg caatcccacc catattgaaa tttgctgaaa tgctttcctc 300  
 aatgtagatc tgtgnaatac accaggggaag ttcanctttt ccggtgt 347

<210> 2494  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2494

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 tggttttgaa ggtttcgaga agangctgga aatatccttt ttccagccgg gactttttgc 120  
 tgacctgag ggaatgggtt taagagctct tgcaaagtcc cagttggatg agatacttac 180  
 accggctgct tgcaccattg tttcatctcn cagaaatgat catgtcgact cctatgttct 240  
 gtctgagtc agtctctttg tttatgcta caagatcatc atcaaaacct gtggtactac 300  
 aaagctactg cttg 314

<210> 2495  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2495

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 atgactgtta attctggcat tagaaaaatt cttccagatt ctgagatttg tgactttgac 120  
 tttgagccat gtggttattc aatgaactct gttgaagggtg ctgctgtttc taccattcat 180  
 gttaccccag aagatgggtt cagttatgca agctttgaga ctgttgggta tgacttcaaa 240  
 gtggtgaatc tgaacgaaat ggtaagagg gtattggcat gttttctccc aactgagttc 300  
 tctgttgcaag ttca 314

<210> 2496  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max

<400> 2496

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tttgtgactt tgactttgag ccatgtgggtt attccaatga actctgttga aggtgctgct 120  
gtttctacca ttcatgttac ccagagaagat ggtttcagtt atgcaagctt tgagactgtt 180  
gggtatgact caaagtgggtg aatctgaacg aaatgggtta gaggggtattg gcatgttttc 240  
tcccaactga gttctctgtt gcagttcatg tggatgggtgc aagcaagttg tttgatcaga 300  
cgtgttttct ggatgttaag 320

<210> 2497

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2497

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tggttttgaa ggtttcgaga aaaggttgga aatatccttt ttccagccgg gactttttgc 120  
tgacctgaa ggaaggggtc taagggctct taaaaatcc cagttgggtg agattctaac 180  
accagctgct tgcaccattg tttcttcgct caaaaacgat aatgtcgact cctatgttct 240  
atctgagtc agcctctttg tttatgccta caagatcatc atcaaaacct gtg 293

<210> 2498

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2498

nnntaanang tcgnangcac gcgtacgtaa gctcngaatt cggctcgagc ttcaaagtgg 60  
tgaatctgaa cgaaatgggtt aagaggggtat tnnctgttt tctcccaact gagttctctg 120  
ttgccgttca tgtgnatggg gcaagcaagt tgtttgatca gacgtgtttt ctggatgtta 180  
agggatactg tcgcgaagag aggagccacg aagggttggt aatgggtgggt tctcttgtct 240  
acaaaaaatt tgccaagact tgtgactgtg gttcacctag atcaactctg aagtgtctgga 300  
aagaggaaga tgaagaagag tagtttt 327

<210> 2499



<211> 315  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2502  
  
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 titgaagggt tcgagaaaag gttggaaata tcctttttcc agccgggact ttttgctgac 120  
 cctgaaggaa ggggtctaag ggctcttaca aaatcccagt tgggtgagat tctaacacca 180  
 gotgcttgca ccattgtttc ttgctcaaa aacgataatg tcgactccta tgttctatct 240  
 gagtccagcc tctttgttta tgcttacaag atcatcatca aaacctgtgg taactactaa 300  
 gotattgctt gcaat 315

<210> 2503  
 <211> 312  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2503  
  
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 cttccagatt ctgagatttg tgactttgac ttgagccat gtggttattc aatgaactct 120  
 gttgaagggt ctgctgtttc taccattcat gttacccag aagatggttt cagttatgaa 180  
 gctttgagac tgttgggtat gacttcaaag tggatgaatct gaacgaaatg gttaagaggg 240  
 tattggcatg ttttctccca actgagttct ctgttgacgt tcatgtggat ggtgcaagca 300  
 agtngtttga tc 312

<210> 2504  
 <211> 440  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2504  
  
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 ttotttgnnt tcaattaggt ttttgctgct ctcttcaaa ctccgtcttt ccgaatctc 120  
 tctttgtgat tgtgttctgt tctgcttct accgcgatag tttctcttct gaagcatggc 180  
 catggcgggt tctgcaattg gttttgaagg ttctgagaag aggctggaaa tatccttttt 240



ccagccggga ctttttgctg accctgaggg aatgggttta agagctcttg caaagtccca 300  
 gttggatgag atacttacac cggctgcttg caccattggt tcatctctca gaaatgatca 360  
 tgtcgactcc tatggctctg ctgaagtcca gtctctttgg ttatgcctac aagatcatta 420  
 tcaaaacctg gggactaca 440

<210> 2505  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2505

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 ttgaagggtt cgagaaaagg ttggaaatat cttttttcca gccgggactt tttgctgacc 120  
 ctgaaggaag gggcttaagg gctcttaca aatcccagtt gggtgagatt ctaacaccag 180  
 ctgcttgca cattgtttct tcgctcaaaa acgataatgt cgactcctat gttctatctg 240  
 agtcacgct ctttgtttat gctacaaga tcatcatcaa aacctgt 287

<210> 2506  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2506

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 accccagaag atggtttcag ttatgcaagc ttcgaaactg ttggttatga cttcaaagcg 120  
 gtgaatctga acgaaatngt tcagagggtta ttggcatggt tcctcccaac tgagttttct 180  
 gttgcagttc atgtggatgg tgcaagcaag tcgtttgagc agacctgctt tctggatggt 240  
 aagggatact gtcgtgaaga gaggagccac gaagggcttg gaatgggtgg ttctgttg 298

<210> 2507  
 <211> 505  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2507

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tgggtggcca anacnaagca cagaactggc atgtctactc tgcttctgca gattctgtaa 120  
 ctcaatgtga caatgtttac actcttgaga tgtgcatgac tggcctggat agagagaaag 180  
 cacaggtttt ctacaaagaa caatctgctt cagctgccat gatgactgnt aattccggca 240  
 ttagaaaaat tcttccaaat tcccagaatt gngactttgn ntttgaacca tgtggntatt 300  
 caatgaactn tgnttgaaag gcncttggtg ttctacgatt catggtancc ccagaagatg 360  
 ggtcanntat tgcaagcttt gnaaactntt gggatgact ttaaagccgg ngaatntgaa 420  
 cccaaaaggn ttaaanggat ttggcatggt tctccaact taantttctg tncaantcat 480  
 tggggaangt gcaagcaagn ntttt 505

<210> 2508  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max

<400> 2508  
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 caatgaactc tgttgaaggc gctgntgttt ctacgattca tgttacccca gaagatgggt 120  
 tcagttatgc aagcttcgaa actgttggtt atgacttcaa agcggatgaat ctgaacgaaa 180  
 tggttcagag ggtattggca tgtttcctcc caactgagtt ttctgttgca gttcatgtgg 240  
 atggtgcaag caagtcgttt gagcagacct gctttctgga tgtaaggga tact 294

<210> 2509  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max

<400> 2509  
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 agatttgtga ctttgacttt gagccatgtg gttattcaat gaactctgtt gaagggtgctg 180  
 ctgtttctac cattcatgtt accccagaag atggtttcag ttatgcaagc tttgagactg 240  
 ttgggtatga cttcaaagtg gtgaatctga acgaaatggt taagaggga ttggca 296

<210> 2510

<211> 254  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2510  
  
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 gagattctaa caccagctgc ttgcaccatt gtttcttcgc tcaaaaacga taatgtcgac 120  
 tectatgttc tatctgagtc cagcctcttt gtttatgcct acaagatcat catcaaaacc 180  
 tgtggtacta ctaagctatt gcttgcaatc ccacccatat tgaagttcgc tgaaatgctt 240  
 tcccttaatg ttaa 254

<210> 2511  
 <211> 299  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2511  
  
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 tcatgttacc ccagaagatg gtttcagtta tgcaagcttc gaaactgttg gttatgactt 120  
 caaagcgggtg aatctgaacg aaatggttca gagggatttg gcatgtttcc tcccaactga 180  
 gttttctgtt gcagttcatg tggatgggtc aagcaagtcg tttgagcaga cctgctttct 240  
 ggatgttaag ggatactgtc gtgaagagag gagccacgaa gggcttgga tgggtgggt 299

<210> 2512  
 <211> 257  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2512  
  
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 aaaacctgtg gtactactaa gctattgctt gcaatccacc catattgaag ttcgctgaaa 120  
 tgctttccct taatgttaag tctgtgaatt acaccagggg aagtttcatt ttccccagt 180  
 ctcagccata tccccatgc aacttttctg aggaagttgc tattcttgat ggctactttg 240  
 gcaaacttgg tgcagga 257

<210> 2513

<211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2513  
  
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 ctaagctatt gcttgcaatc ccacccatat tgaagtgcg tgaaatgctt tcccttaatg 180  
 ttaagtctgt gaattacacc aggggaagtt tcattttccc cagtgcgcag ccataatcccc 240  
 atcgcaactt ttctgaggaa gttgctattc ttgatggcta ctttggcaaa cttgggtgcag 300  
 gaagcaatgc 310

<210> 2514  
 <211> 322  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2514  
  
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 ctcccaactg agttttctgt tgcagttcat gtggatgggt caagcaagtc gtttgagcag 120  
 acctgctttc tggatgttaa gggatactgt cgtgaagaga ggagccaaga agggcttggg 180  
 atgggtggtt ctgttgtcta ccaaaaattc gggaagacta gtgactgtgg ttcacctaga 240  
 tcaactctaa agtgctggaa tgaggaagat gaggaagagt agtttcctta agtgtcttta 300  
 ttattctgtc cttgtgaaaa ta 322

<210> 2515  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2515  
  
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 ttacaaaatc ccagttgggt gagattctna caccagctgc ttgcaccatt gtttcttcgc 120  
 tcaaaaacga taatgtcgac tcctntgttc tatctganc cagcctcttt gtntatgcct 180  
 acnagatcat catcaaaacc tgtgggtacta ctaagctatt gcttgcnacc ccacccatat 240

tgaagttngc tganatgctt tcccttaatg ttaagtctgt gaattacacc aggggaagtt 300  
tcattttccc cagt 314

<210> 2516  
<211> 283  
<212> nucleic acid  
<213> Glycine max  
<400> 2516

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atggtttcag ttatgcaagc ttgagactg ttgggtatga cttcaaagtg gtgaatctga 180  
acgaaatggg taagagggtg ttggcatgtt ttctcccaac tgagttctct gttgcagttc 240  
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<210> 2517  
<211> 247  
<212> nucleic acid  
<213> Glycine max  
<400> 2517

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attgottgca atcccaccca tattgaagtt cgctgaaatg ctttccctta atgttaagtc 120  
tgtgaattac accaggggaa gtttcatttt cccagtgct cagccatata cccatcgcaa 180  
cttttotgag gaagttgcta ttcttgatgg ctactttggc aaacttggtg caggaagcaa 240  
tgottat 247

<210> 2518  
<211> 336  
<212> nucleic acid  
<213> Glycine max  
<400> 2518

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tttctgagga agttgctatt cttgatggct actttggcaa acttggtgca ggaagcaatg 120  
cttatatattt ggggtggccaa gacaaagcac agaactggca tgtctactct gcttctgcag 180

attctgtaac tcaatgtgac aatgtttaca ctcttgagat gtgcatgact ggcttgata 240  
gagagaaagc acaggttttc tacaagaac aatctgcttc agctgccatg atgactgtta 300  
attccggcat tagaaaaatt cttccagatt ccgaga 336

<210> 2519  
<211> 306  
<212> nucleic acid  
<213> Glycine max  
<400> 2519

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ctaagctatt gcttgcaatc ccacccatat tgaagttcgc tgaaatgctt tcccttaatg 120  
ttaagtctgt gaattacacc aggggaagtt tcattttccc cagtgtcag ccatatcccc 180  
atcgcaactt ttctgaggaa gttgctattc ttgatggcta ctttggcaaa cttggtgcag 240  
gaagcaatgc ttatatatttg ggtggccaag acaaagcaca gaactggcat gtctactctg 300  
cttctg 306

<210> 2520  
<211> 247  
<212> nucleic acid  
<213> Glycine max  
<400> 2520

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gaaatgtttc cctcaatggt agatctgtga attacaccag gggaagtttc atctttcccg 180  
gtgtcagcc ctatcccat cgcaactttt ctgaggaagt tgctattctt gatggctact 240  
ttggcaa 247

<210> 2521  
<211> 282  
<212> nucleic acid  
<213> Glycine max  
<400> 2521

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gcatgactgg cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag 120  
 ctgccatgat gactgttaat tccggcatta gaaaaattct tccagattcc gagatttgtg 180  
 actttgactt tgnaccatgt ggttattcaa tgaactctgt tgaaggcgct gctgtttcta 240  
 cgattcatgt taccacagaa gatggtttca gttatgcaag ct 282

<210> 2522  
 <211> 305  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2522

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 tattgcttgc aatcccaccc atattgaagt tcgctgaaat gctttccctt aatgttaagt 120  
 ctgtgaatta caccagggga agtttcattt tccccagtcg tcagccatat ccccatcgca 180  
 acttttntga ggaagttgct attcttgatg gctactttgg caaacttggt gcaggaagca 240  
 atgcttatat tttgggtggc caagacaaag cacagaactg gcatgtctac tctgcttctg 300  
 cagat 305

<210> 2523  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2523

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 cgggactttt tgctgaccct gagggaatgg gttaagagc tcttgcaaag tcccagttgg 180  
 atgagatact tacaccggct gcttgacca ttgtttcatc tctcagaaat gatcatgtcg 240  
 actcctatgt tctgtctgag tccagtctct tgtttatgcc tacaaga 287

<210> 2524  
 <211> 276  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2524

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ggtgctgctg tttctacat tcatgttacc ccagaagatg gtttcagtta tgcaagcttt 180  
gagactgttg gncatgactt caaagtggg aatctgaacg aaatgggttaa gaggggtattg 240  
gcatgttttc tcccaactga gttcgtgttg cagttc 276

<210> 2525  
<211> 302  
<212> nucleic acid  
<213> Glycine max  
<400> 2525

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gatactgtcg cgaagagagg agccacgaag ggcttggaat ggggtggttct cttgtctacc 180  
aaaaatttgc caagacttgt gactgtggtt cacctagatc aactctgaag tgntggaaag 240  
aggaagatga agaagagtag ttttcttaag tgtctttatt atgtccttgc gaaaataagt 300  
cc 302

<210> 2526  
<211> 274  
<212> nucleic acid  
<213> Glycine max  
<400> 2526

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ctactaagct attgcttgca atcccacca tattgaagtt cgctgaaatg ctttccctta 180  
atgttaagtc tgtgaattac accaggggaa gtttcatttt cccagtgct cagccatatc 240  
cccatcgcaa cttttctgag gaagttgcta ttct 274

<210> 2527  
<211> 264  
<212> nucleic acid  
<213> Glycine max



<400> 2527

ctgcaattgg ttttgaaggt ttcgagaagn ngctggaaat atcctttttc cagccgggac 60  
tttttgctga ccctgagggg atgggtttta gagctcttgc aaagtcccag ttggatgaga 120  
tacttacacc ggctgcttgc accatangtt tcatctctca gaaatgatca tgtcgactcc 180  
tatgttctgt ctgagtnacg tctctttgtt tatgcctaca agatcatcat caaaacctgt 240  
ggtatacaaa gctactgttg cant 264

<210> 2528

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2528

ntgcacgcgt acgtaagctc ggaattcggc tcgaggctac tgcttgcaat cccacccata 60  
ttgaaatttg ctgaaatgct ttccctcaat gttagatctg tgaattacac caggggaagt 120  
ttcatctttc ccgggtgctca gccctatccc catcgcaact tttctgagga agttgctatt 180  
cttgatggct actttggcaa gcttagtgca ggaagcaatg ottatatttt ggggtggccaa 240  
gacaaatcac agaactggca tgtctactct gcttctgcag attctgtaa 289

<210> 2529

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2529

tcnnangcac gcgtacgtna gctcggaatt cggctcgagg cttgcaatcc cacccatatt 60  
gaagttcgct gaaatgcttt cccttaatgt taagtctgtg aattacacca ggggaagttt 120  
cattttcccc agtgctcagc catatcccca tcgcaacttt tctgaggaag ttgctattct 180  
tgatggctac tttggcaaac ttggtgcagg aagcaatgct tatatttttg gtggccaaga 240  
caaagcacag aactggcatg tctactctgc ttctgcagat tctgtaactc aatgtgacaa 300  
tgtttacatc t 311

<210> 2530

<211> 308

<212> nucleic acid

<213> Glycine max

<400> 2530

tntangcac gcgtagctna gctcggaatt cggctcgagc aggaagcaat gcttatatnt 60  
tgggtggcca agacaaatca cagaactggc atgtctactc tgcttctgca gattctgtaa 120  
ctccatgoga caatgtttac actctagaga tgtgcatgac tggcctggat agagagaaaag 180  
cacaggtttt ctacaaagaa caatctgctt cagctgccat tatgactgtt aattctggca 240  
ttagaaaaat tcttcagat tctgagattt gtgactttga ctttgagcca tgtggttatt 300  
caatgaac 308

<210> 2531

<211> 292

<212> nucleic acid

<213> Glycine max

<400> 2531

tgcangcac gcgtagctaa gctcggaatt cggctcgagc aaagcacaga actggcatgt 60  
ctactctgct tctgcagatt ctgtaactca atgtgacaat gtttactc ttgagatgtg 120  
catgactggc ctggatagag agaaagcaca ggttttctac aaagaacaat ctgcttcagc 180  
tgccatgatg actgttaatt cgggcattag aaaaattctt ccagattccg agatttgtga 240  
ctttgacttt gaaccatgtg gttattcaat gaactctgtt gaaggcgtg ct 292

<210> 2532

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 2532

cnaangcacg cgtacgtaag ctcggaattc ggctcgaggc agattctgta actcaatgtg 60  
acaatgttta cactcttgag atgtgcatga ctggcctgga tagagagaaa gcacaggttt 120  
tctacaaaga acaatctgct tcagctgcc tgaatgactgt taattccggc attagaaaaa 180  
ttcttcagga ttccgagatt tgtgactttg actttgaacc atgtggttat tcaatgaact 240  
ctggtgaagg cgctgctgtt tctacgattc atgttactcc agaag 285

<210> 2533

<211> 326  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2533  
  
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 gatctgtctt ctactgcgat agtttctcta ctgttacatg gccatggcgg tttccgcaat 120  
 tggttttgaa ggtttctgaga aaaggttgga aatatccttt ttccagccgg gactttttgc 180  
 tgacctgaa ggaaggggtc taagggctct taaaaatcc cagttgggtg agatctaaca 240  
 ccagctgctt gcaccattgt ttcttcgctc aaaaaacgat aatgtcgact cctatgttct 300  
 atctgagtcc agcctctttg tttatg 326

<210> 2534  
 <211> 502  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2534  
  
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 gntttcaatt aggttggttt gntgctctcc ttcaaactcc atctttccaa atcctctctt 120  
 tgcgattgng ttttgatctg ctctctactg cgatagnttc tctactgnta catggccatg 180  
 gcggtttccg caattgggtt tgaaggnttc gagaaaaggn tggaaatatc ctttttccaa 240  
 ccgggacttt ttgctgacct tgaaagaagg ggtctaaang gctnttacao aatccaagtg 300  
 ggtgagattc taacaccagc tgnttgnacc attgggttctt ngctnaaaaa cgatnatgnc 360  
 cacttctatg gtctatctna gttcangctt tttgggtatg cctaccaaga tcattattna 420  
 aaactnnggg accacctaac tattgggttn aatcccccnt atttgaaatt gcttnaanng 480  
 ctttccctta aggttaaact gg 502

<210> 2535  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2535  
  
 nangtcgcan gcacgcgtac gtaagctcgg aattcggctc gaggctactg cttgcaatcc 60

cacccatatt gaaatttgct gaaatgcttt ccctcaatgt tagatctgtg aattacacca 120  
 ggggaagttt catctttccc ggtgctcagc cctatcncca tcgcaacttt tctgaggaag 180  
 ttgctattct tgatggctac tttggcaagc ttagtgcagg aagcaatgct tatattttgg 240  
 gtggccaaga caaatcacag aactggcatg tctactctgc ttctgcagat t 291

<210> 2536  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2536

gcangcagc gtacgtaagc tcgggaattc ggctcgagng ccaagacaaa gcacagaact 60  
 gggcatgtct actctgcttc tgcagattct gtaactcaat gtgacaatgt ttacactctt 120  
 gagatgtgca tgactggcct ggatagagag aaagcacagg tttttacaa agaacaatct 180  
 gcttcagctg ccatgatgac tgttaattcc ggcattagaa aaattcttcc agattccgag 240  
 atttgtgact ttgactttga accatgtggg tattcaatga actctgttga aggcgctgct 300  
 gttttctac 308

<210> 2537  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2537

acgtcgcang cacgcgtacg taagctcgng aattcggctc gagggcaagc ttagtgcagg 60  
 aagcaatgct tatattttgg gtggccaaga caaatcacag aactggcatg tctactctgc 120  
 ttctgcagat tctgtaactc catgcgacaa tgtttacact ctagagatgt gcatgactgg 180  
 cctggataga gagaaagcac aggttttcta caaagaacaa tctgcttcag ctgccattat 240  
 gactgttaat tctggcatta gaaaaattct tccagattct gagatttgtg actttgactt 300  
 tgagccat 308

<210> 2538  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max

<400> 2538

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gtactacaaa gctactgctt 60

gcaatcccac ccatattgaa atttgctgaa atgctttccc tcaatgttag atctgtgaat 120

tacaccaggg gaagtttcat ctttcccggt gtcagccct atccccatcg caacttttct 180

gaggaagttg ctattcttga tggctacttt ggcaagetta gtgcaggaag caatgcttat 240

attttgggtg gccaaagaaa atcacagaac tggcatgtct a 281

<210> 2539

<211> 299

<212> nucleic acid

<213> Glycine max

<400> 2539

antcgcatgc acgcgtacgt nagctcgga ttcggctcga gattctggca ttagaaaaat 60

tottccagat tctgagattt gtgactttga ctttgagcca tgtggttatt caatgaactc 120

tgttgaaggt gctgctgttt ctaccattca tgttacccca gaagatgggt tcagttatgc 180

aagctttgag actgtttgggt atgacttcaa agtgggtgaat ctgaacgaaa tggttaagag 240

gggatgggca gttttcttcc caatgagttc tctgttgag ttcagtggga ggtgcaaca 299

<210> 2540

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2540

ntgnncgct acgtgagtc ggaattnggc tcgagctcga gccgttcgag aagaggctgg 60

aaatatcctt tttccagcng gnactttttg ctgaccctga ggncatgggt tnangagntc 120

ttgcaaagtc ccagttggat gagatannta nacncgctgc ttgcaccatt gtttcatctc 180

tcagaaatga tcatgncgan tcctatgtnc tgtctgagtc cagtntcttn gtntatgcct 240

acaagatcat catcaaaacc tgnngtacta caaagctact gctt 284

<210> 2541

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 2541

gcgtacgtaa gctcggaatt cggctcgagn ttaggttttt ttgttgctct ccttcaaact 60  
ccatctttcc aaatcctctc ttgcgattg tgttttgatc tgcttctac tgcgatagtt 120  
tctctactgt tacatggcca tggcggtttc cgcaattggt tttgaagggt tcgagaaaag 180  
gttggaata tcctttttcc agccgggact ttttgctgac cctgaaggaa ggggtctaag 240  
ggctcttaca aaatcccagt tgggtgagat tctaacacca gctgcttgca ccatggt 297

<210> 2542

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 2542

tcgnngcacg cgtacgtaag ctcggaattc ggctcgaggg ccaagacaaa tcacagaact 60  
ggcatgtcta ctctgcttct gcagattctg taactccatg cgacaatggt tacactctag 120  
agatgtgcat gactggcctg gatagagaga aagcacaggt tttctacaaa gaacaatctg 180  
cttcagctgc cattatgact gttaattctg gcattagaaa aattcttcca gattctgaga 240  
tttgtgactt tgacttgagc catgtgggta ttcaatgaac tctgttgaag gtgctgct 298

<210> 2543

<211> 390

<212> nucleic acid

<213> Glycine max

<400> 2543

ttctgatat ccatccagat tgatagttca ttgcatgcaa ccatagtttt attaggtttt 60  
ttcttctttg ttttcaatta ggtttttgct gctctccttc aaactccgtc tttccgaatc 120  
ctctctttgt gattgtgttc tgttctgctt cctaccgca tagtttctct tctgaagcat 180  
ggccatggcg gtttctgcaa ttggttttga aggtttcgag aagaggctgg aaatatcctt 240  
tttccagccg ggactttttg ctgaccctga gggaatgggt ttaagagctc ttgcaaagtc 300  
ccagttggat gagatactta caccggctgc ttgcaccatt ggttcatctc tcagaaatga 360  
tcatgtcgac ttctaaggtc tggtgaanc 390

<210> 2544

<211> 284  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2544  
  
 nncngcacgc nnacgtaagc tcggaattcg gctcgagggg aagtttcatt ttccccagtg 60  
 ctcagccata tccccatcgc aacttttctg aggaagttgc tattcttgat ggctactttg 120  
 gcaaacttgg tgcaggaagc aatgcttata ttttgggtgg ccaagacaaa gcacagaact 180  
 ggcatgtcta ctctgcttct gcagattctg taactcaatg tgacaatgtt tacactcttg 240  
 agatgtgcat gactggcctg gatagagaga aagcacaggt tttc 284

<210> 2545  
 <211> 295  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2545  
  
 gtgcnnngca cgcgtacgta agctcggaat tcggctcgag ctcagccata tccccatcgc 60  
 aacttttctg aggaagttgc tattcttgat ggctactttg gcaaacttgg tgcaggaagc 120  
 aatgcttata ttttgggtgg ccaagacaaa gcacagaact ggcatgtnta ctctgcttct 180  
 gcagattctg taactcaatg tgacaatgtt tacactcttg agatgtgcat gactggcctg 240  
 gatagagaga aagcacaggt tttctacaaa gaacaatctg cttcagctgc catga 295

<210> 2546  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2546  
  
 ngtcgcatgc acgcgtacgt aagctcgga ttcggctcga gtgctcagcc atatccccat 60  
 cgcaactttt ctgaggaagt tgctattctt gatggctact ttggcaaact tgggtgcagga 120  
 agcaatgctt atattttggg tggccaagac aaagcacaga actggcatgt ctactctgct 180  
 tctgcagatt ctgtaactca atgtgacaat gtttacctc ttgagatgtg catgactggc 240  
 ctggatagag agaaagcaca ggttttctac aaagaacaat ctgcttcagc tggccatgat 300  
 gactgttaat 310

<210> 2547  
 <211> 374  
 <212> nucleic acid  
 <213> Glycine max

<400> 2547

cacttaaant cgatgcacgc gtacgtaagc tcggaattcg gctcgagctt gcaccattgt 60  
 ttcatctctc agaaatgata atgtcgacnc ctatgttctg tctgagtcca gtctctttgt 120  
 ttatggctta caagatcata atcaaacct gtggtactac aaagctactg cttgcaatcc 180  
 caccatatt gaaatttgct gaaatgcttt ccctcaatgt taatctgtga attacaccag 240  
 gggaagtttc atctttcccg gtgtcagcc ctatcccat cgcaactttt ctgaggaagt 300  
 tgctattctt gatggctatt tggcaagctt agtgcaggaa gcaagcttat atttgggtgg 360  
 ccagacaaat caca 374

<210> 2548  
 <211> 343  
 <212> nucleic acid  
 <213> Glycine max

<400> 2548

nnntannccc canacgtcgc angcacgcnt acgtnacgtc ggaattcggc tcgagntctt 60  
 ctttggtttc aatnaggttt tttgtgtgct ctcttcaaa ctccatcttt ccaaatectc 120  
 tctttgcgat tgtgttttga tctgttctt actgcgatag tttctctact gttacatggc 180  
 catggcggtt tccgcaattg gttttgaagg ttctgagaaa aggttggaaa tctcttttt 240  
 ccagccggga ctttttgctg accctgaagg aaggggtcta agggctctta caaatccca 300  
 gttgggtgag attctaacac cagctgcttg caccattggt tct 343

<210> 2549  
 <211> 292  
 <212> nucleic acid  
 <213> Glycine max

<400> 2549

nnnnnggcng acgtcgcang cacgcgtacg taagctcgga attcggctcg agcttacacc 60  
 ggctgcttgc accattgtnt catctctcag aatgaacat gtcgactcct atgtgctgtc 120



tgagtnacgt ctctntgttt atgcctacaa gatcatcatc aaaacctgtg gtactanaaa 180  
gctactgctt gcaatcccan ccatattgan atntgctgna atgctttccc ncaatgnag 240  
atctngaat tacaccaggg gaagtttctt cttncctggg gctcagccct at 292

<210> 2550  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2550

ngtcgcatgc acgontacgt aagcncggga attcggctcg agcaaatcc cagttgggtg 60  
agattctaac accagctgct tgcaccattg tntcttcgct caaaaacgat aatgtcgact 120  
cctatgtncct atctgagtec agcctctttg tttatgccta caagatcatn atcaaaacct 180  
gtggactac taagctattg cttgcaatcc cacccatatt gaagttcgct gaaatgcttt 240  
ccctaagtgt aagtctgtga attacaccag gggaagtttc atttccccgt gctcagccat 300

<210> 2551  
<211> 291  
<212> nucleic acid  
<213> Glycine max  
<400> 2551

nngtcgcatg cacgcgtacg taagctcgga attcggctcg aggttcatgt ggatgggtgca 60  
agcaagtcgt ttgagcagac ctgctttctg gatgttaagg gatactgtcg tgaagagagg 120  
agccacgaag ggcttggaat ggggtggttct gttgtctacc aaaaattcgg gaagactagt 180  
gactgtgggt cacctagatc aactctaaag tgctggaatg aggaagatga ggaagagtag 240  
tttctttaag tgtctttatt attctgtcct gtgaaaataa gtctgggttt c 291

<210> 2552  
<211> 294  
<212> nucleic acid  
<213> Glycine max  
<400> 2552

acgtcgcatg cacgcgtacg taagctcgga attcggctcg agtccaaatc ctctctttgc 60  
gattgtgttt tgatctgctt cctactggcg atagtttctc tactgttaca tggccatggc 120

ggtttccgca attggttttg aaggtttcga gaaaagggtg gaaatatcct ttttccagcc 180  
 gggacttttt gctgaccctg aaggaagggg tctaagggtt cttacaaaat cccagttggg 240  
 tgagattcta acaccagctg cttgcaccat tgtttcttcg ctcaaaaacg ataa 294

<210> 2553  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2553

nangnacgg tncgtnacgt cacgtatanc tcggcattcg gctcgagctt tgttttcaat 60  
 taggtttttt tgttgctctc cttcaaactc catctttcca aatcctctct ttgcgattgt 120  
 gttttgatct gcttctact gcgatatgtt ctctactgtt acatggccat ggcgggtttcc 180  
 gcaattgggt ttgaagggtt cgagaaaagg ttggaaatat cttttttcca gccgggactt 240  
 tttgctgacc ctgaaggaag ggggtctaagg gctcttaca aatcccagtt gggtgaga 298

<210> 2554  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2554

acgcgtacgt aagctcggaa ttcggctcga gggtgcaagc aagtcgtttg agcagacctg 60  
 ctttctggat gttaagggtt actgtcgtga agagaggagc cacgaagggc ttggaatggg 120  
 tggttctgtt gtctacaaa aattcgggaa gactagtac tggtggttcac ctagatcaac 180  
 tctaaagtgc tggaatgagg aagatgagga agagtagttt ccttaagtgt cttattattc 240  
 tgtccttggtg aaaataagtc tggttttcca gata 274

<210> 2555  
 <211> 263  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2555

ncgtcgcatt cacgcgtacg taagctcggg attcggctcg agggtgcaag caagtcgttt 60  
 gagcagacct gctttctgga tgtaaggga tactgtcgtg aagagaggag ccacgaaggg 120

cttggaatgg gtggttctgt tgtctaccaa aaattcggga agactagtga ctgtggttca 180  
 cctagatcaa ctctaaagtg ctggaatgag gaagatgagg aagagtagtt tccttaagtg 240  
 tctttattat tctgtccttg tga 263

<210> 2556  
 <211> 275  
 <212> nucleic acid  
 <213> Glycine max

<400> 2556

cacgcgtacg taagctcgga attcggctcg aggggtgcaag caagtcgttt gagcagacct 60  
 gctttctgga tgttaagggg tactgtcgtg aagagaggag ccacgaaggg cttggaatgg 120  
 gtggttctgt tgtctaccaa aaattcggga agactagtga ctgtggttca cctagatcaa 180  
 ctctaaagtg ctggaatgag gaagatgagg aagagtagtt tcctaaagtgt cttattattc 240  
 tgtcctgtga aaataagtct ggttttccag atacg 275

<210> 2557  
 <211> 280  
 <212> nucleic acid  
 <213> Glycine max

<400> 2557

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gaagcaatgc ttatatatttg 60  
 ggtggccaag acaaatcaca gaactggcat gtctactctg cttctgcaga ttctgtaact 120  
 ccattgcgaca atgtttacac tctagagatg tgcattgact gcttgatag agagaaagca 180  
 caggttttct acaaagaaca atctgcttca gctgccatta tgactgttaa ttctggcatt 240  
 agaaaaattc ttccagattc tgagatttgt gactttgact 280

<210> 2558  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max

<400> 2558

gtcgcangcg tacgtaagct cggaattcgg ctcgagccca gttgggtgag attctaacac 60  
 cagctgcttg caccattgtt tcttcgtcga naaacgataa tgtcgactcc tatgtttctat 120

ctgagtccag cctctttgtt tatgcctaca agatcatcat caaaacctgt ggtactacta 180  
agctattgct tccaatccca cccatattga agttcgctga aatgcttncc ttaatgttaa 240  
gtctgtgaat acaccagggg aagtttcatt tccccagtgc tcagccatat ccccatcgca 300  
atcttctgan g 311

<210> 2559  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
<400> 2559

ctnecatgac gtaagtnago tcggaattcg gctcgagggt tttttgttg ctctccttca 60  
aactccatct ttcctaatcc tctctttgcn ttgtgttttg atctgcttcc tactgcgata 120  
gtttctctac tgttacatgg ccattggcggg ttccgcaatt ggttttgaag gtttcgagaa 180  
aagggttgaa atctcctttt tccagccggg actttttgct gaccctgaag gaaggggtct 240  
aagggtctct acaaaatccc agttgggtga gattctaaca ccagctgctt gc 292

<210> 2560  
<211> 288  
<212> nucleic acid  
<213> Glycine max  
<400> 2560

ggntatgcac ncnacgtgn gcncngagtt cggcncgngg ggggnttccg naatnggttt 60  
tgaagggttc gagaaaaggt tggaaatct cttntccan nccgggactt tntgctgacc 120  
ctggnaggaa ggggtctaca gggctcttac aaaatcccag ttgggtgaga ttctaacacc 180  
agctgcttgc accatgtttc ttncntcana aacgatnntg tcgactccta tgttctatct 240  
gagtnacgcc tctttgttna tgctacaag atcatcancn anacctgt 288

<210> 2561  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2561

tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc ncantngggg gagattcnaa 60

naccagctgc ttgcacnant gtttcttcgc tcaaaaacna gcaatgtcga ctcctatggt 120  
ctatctgagt ccagcctctt tgtttatgcc tncaaganca tcatcaaaac ctgggtacta 180  
ctaagctatt gcttccaatc ccacccatat tgaagttcgc tgaaatgctt tcccttaatg 240  
ttaagtctgt gaattacacc aggggaagtt tcattttccc cagtgtcag ccatatcccc 300

<210> 2562  
<211> 236  
<212> nucleic acid  
<213> Glycine max  
<400> 2562

ncgcgtacgt aagctcggaa ttccggtcga gcagcattcg ttgcggcatt ttaatcgatt 60  
tatccaagca ggactgaatg aactaatgga gtctaaacnc gggaaaaaga agtctagtag 120  
tagtagtagt aaatcattgt tctacgaagc tcccctcgga tacagcattg aagacgttag 180  
accaaacggt ggaatcaaga aattcagatc tgctgcttac tccaactgcg ctcgcn 236

<210> 2563  
<211> 285  
<212> nucleic acid  
<213> Glycine max  
<400> 2563

ncncgtacgt aagctcggaa ttnggctcga gctcgagncg gttcatgtgg ntngtgcaag 60  
caatttgtnt gatcanacgt gnnttctgga tgtaagga tactgtcgcg aagagaggag 120  
ccacgaaggg cttggaatgg gtggttctct tnnctacca aaaatttgcc aagacttggt 180  
actgtgcgct cacctagatc aactctgaag tgctggaaag aggaagatga agaagagtag 240  
ttttcttaag tgtctttatt atgtccttgc gaaaataagt ccggt 285

<210> 2564  
<211> 286  
<212> nucleic acid  
<213> Glycine max  
<400> 2564

acgtcgcgtg cagcgtacg taagctcgga attcggctcg agaacttttc tgaggaagtt 60  
gctattcttg atggctactt tggcaagctt agtgcaggaa gcaatgctta tattttgggt 120

ggccaagaca aatcacagaa ctggcatgtc tactctgctt ctgcagattc tgtaactcca 180  
 tgcgacaatg ttacactct agagatgtgc atgactggcc tggatagaga gaaagcacag 240  
 gttttctaca aagaacaatc tgcttcagct gccattatga ctgttg 286

<210> 2565  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max

<400> 2565

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ngttttgggtt gattcaaggc 60  
 cttcacagca ttcgntgagg ctttttnatn gatntntcca agcaggactg natgaactaa 120  
 tggagtctaa aggtgggaaa aagaagtcta gtagtagtag tagtanatca ttttctacga 180  
 agtcccccgc ggntacagca ttgaagangt tagaccaaac ggtggaatca agannttcag 240  
 atctgctgct tactccaact gcgctcgcaa accttctgn tatccatcca gattga 296

<210> 2566  
 <211> 492  
 <212> nucleic acid  
 <213> Glycine max

<400> 2566

ccacgcgtcc gcttcaaadc acacactctc ttcaatttct agggttttgc tattgctttg 60  
 cctccgttcc ccngntctca caaaaacaac gccttttctc ttctccttgc tatctattct 120  
 ttgcgtttgg tttttggttg attgaaggca ttcacagcng taattcggtg ctgcatttta 180  
 atcgatttat ctaaccagga ctgaatgacc taatggagtc taaagggtggg aaaaagaagt 240  
 ctagtagtag tagtagtaaa tcaatttttt acgaagctcc cctcggatac agcattgaag 300  
 acgttagacc aaacggtgga atcaagaaat tcagatctgc tgcttactct aactgttctc 360  
 gcaaaccatc ctgatacaca tccggattga tagttcgttg catgcaacca tagttttatt 420  
 aggatttttt cttctttgnt ttcaattagg tttttttggt gctctcttcc aaactccatc 480  
 tttccaaatc ct 492

<210> 2567  
 <211> 298  
 <212> nucleic acid

<213> Glycine max  
 <400> 2567  
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 ggtttttggg tgattgaagg cattcacagc gtaattcggt gctgcatttt aatcgattta 120  
 tctaaccagg actgaatgac ctaatggagt ctaaagggtg gaaaaagaag tctagtagta 180  
 gtagtagtaa atcaattttt tacgaagctc ccctcgata cagcattgaa gacgttagac 240  
 caaacgggtg aatcaagaaa ttcagatctg ctgcttactc taactgttct cgcaaacc 298

<210> 2568  
 <211> 277  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2568  
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 aganaagggt ggaaanctcc tttannccag ccgggacttt anggetgaen ctgaaggaag 120  
 aggtctaagg gctcttacia aatcccagtt gggtagatt ctaacaccag ctgcttgac 180  
 cattgtttct tcgctcaaaa acgataatgt cgactcctat gttctatctg agtcagcct 240  
 cttgtttatg cctacnagat catcatcana acctgtg 277

<210> 2569  
 <211> 307  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2569  
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 tgtctgctct ctttcaaact ccgtctttcc gaatcctctc tttgtgattg tnttctgttc 120  
 tgcttcttac cgcgatagtt tctcttctga agcatggcca tggcggtttc tgcaattggg 180  
 tttgaagggt tcgagaagag gctggaaata tcctttttcc agccgggact ttttgctgac 240  
 cctgagggna tgggttttag agctcttgn aagtcccagt tggntgagat acttacaccg 300  
 gctgctt 307

<210> 2570

<211> 245  
<212> nucleic acid  
<213> Glycine max

<400> 2570

ggtttttggct gctctccttc aaactccgtc tttccgaatc ctctctttgt gatttgtgttc 60  
tgttctgctt cctaccgcga tagtttctct tctgaagcat ggccatggcg gtttctgcaa 120  
ttggttttga aggtttcgag aagaggctgg aaatatcctt tttccagccg ggactttttg 180  
ctgaccctga gggaatgggt ttaagagctc ttgcaaagtc ccagttggat gagatactac 240  
accgg 245

<210> 2571  
<211> 326  
<212> nucleic acid  
<213> Glycine max

<400> 2571

nnncgcangc acgcgtacgt aagctcggaa ttcggctcga ggttttctcc caactgagtt 60  
ctctgtttgca gttcatgtgg atggtgcaag caagttgttt gatcagacgt gttttctgga 120  
tggttaaggga tactgtcgcg aagagaggag ccacgaaggg cttggaatgg gtggttctct 180  
tgtotaccaa aatttgccaa gacttgtgac tgtggttcac ctagatcaac tctgaagtgc 240  
tggnaagagg nagatgcaga agagtagttt tcttaagtgt ctttattatg tccttgcgaa 300  
aataagtccg gttttccaga cagtga 326

<210> 2572  
<211> 281  
<212> nucleic acid  
<213> Glycine max

<400> 2572

agttgcgcgn acncgtacgt aagctcggaa ttcggctcga gaaatcattg ttctangaag 60  
ctccccctcgn atanagcatn ggngacgtta gaccaannng tggaatcaag aaattcagat 120  
ctgntgctta ctccaactgc gctcgcaaan ccttcctgat atccatcng attgatagtt 180  
cattgcatgc aaccatagtt tnattaggtt ttntcttctt tgttttcaat taggtttttg 240  
ctgctctcct tcaaactccg tctttccgaa tcctctcttt g 281



<210> 2573  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2573  
  
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 aattaggttt ttgctgctct ccttcaaact ccgtctttcc gaatcctctc tttgtgattg 120  
 tgtttctgttc tgcttcttac cgcgatagtt tctcttctga agcatggcca tggcgggtttc 180  
 tgcaattggt tttgaaggtt tcgagaagag gctggaaata tcctttttcc agccgggact 240  
 ttttgctgac cctgagggaa tgggtttaag agctcttgca aagtcccagt tggatgag 298

<210> 2574  
 <211> 450  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2574  
  
 aagggggatt ctgtangngn natngganng gacaacatan aagcgtatg acgtcgcatg 60  
 cacgcgtacg taagctcgga attcggctcg agtttatgcc tacaagatca tcatcaaaac 120  
 ctgtggtacn nntaagctat tgcttgcaat cccaccata ttgaagttcg ctgaaatgct 180  
 ttcccttaat gttaagtctg tgaattacac caggggaagt ntcattttcc ccagtgtcga 240  
 nccatatcnn catcgcaagn tttntgagga agattnnant gttnttngtt antntnnn 300  
 ncttttttna tttacttaac ttatnatttg nncnttttat ntttaagcat natnactnna 360  
 nntttttnag gnggggtgtn ttntttnttn ntctttnttn tttttttnnn attcanttta 420  
 ttngttnttn tntntntnnn ntnttcnttt 450

<210> 2575  
 <211> 218  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2575  
  
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 ggtttcgaga aaaggttgga aatatccttt ttccagccgg gactttttgc tgaccctgaa 120

ggaaggggtc taagggctct tacaaaatcc cagttgggtg agatnctaac accanctgct 180  
 tgnancattg tntcttcgct caaaaacgat aatgtcgn 218

<210> 2576  
 <211> 428  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2576

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 caagacttgt gactgngggt cacctanato aactctgaag tgctggaaag aggaagatga 180  
 anaanagtat ttttcttaag tgtctttatt atgtccttgc naaaataagt ccngttttnc 240  
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 gtttggaact tttattgntc tactattacc atttgaacat cgnatggatt ttaataaaan 360  
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 naattctn 428

<210> 2577  
 <211> 312  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2577

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 ctaagctatn gcttgcaatc ccacccatat tgnagtctgc tgaaatgctt cccttaatgt 180  
 naagtctgtg aattacacca ggggaagnnt cattttcccc agtgcncagc canatncna 240  
 togcaanttt tnngaggnag tcnccattcc tggatngcct actttggcaa acttggtgog 300  
 gangcaatgc tt 312

<210> 2578  
 <211> 261  
 <212> nucleic acid  
 <213> Glycine max

<400> 2578

gtcgcacgca gcgtacgtaa gctcggaatt cggctcgagc agccatatcc ccacgcaac 60

ttttctgagg aagttgctat tcttgatggc tactttggca aacttggtgc aggaagcaat 120

gcttatatatt tgggtggcca agacaaagca canaactggc atgtctactc tggtttctgc 180

agattctgaa ctcaatgtgc caatgnttac actcntgagn tgnngcatga ctggctggat 240

agagagaang cncaggtttt c 261

<210> 2579

<211> 279

<212> nucleic acid

<213> Glycine max

<400> 2579

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ctagtagtag tagtagtaaa tcaatttttt acgaagctcc cctcggttac agcattgaag 120

acgttagacc aaacgggtgga atcaagaaat tcagatctgc tgcttactct aactgttctc 180

gcaaaccatc ctgatacaca tccggtattga tagttcgttg catgcaacca tagttttatt 240

aggatttttc tcttgttttc aattagggtt tttgtgtgc 279

<210> 2580

<211> 234

<212> nucleic acid

<213> Glycine max

<400> 2580

tcgangcacg cgtacgtaag ctcggaattc ggctcgagaa gcaatgctta tattttgggt 60

ggccaagaca aagcacagaa ctggcatgtc tactctgctt ctgcagattc tgtaactcan 120

tgtgacaatg ttacactct tgagatgtgc atnactggcc tggatagaga gaaagcacag 180

gttttcnaca aagaacaatc tgcttcagct gccatgatga ctgnaannc cggc 234

<210> 2581

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 2581

gcacgcacgc gtacgtaagc tcggaattcg gctcgagntt cttctttgtt ttcaattagg 60  
 tttttttgtt gctctccttc aaactccatc tttccaaatc ctctctttgc gatttgtgtt 120  
 tgatctgctt cctactgcga tagtttctct actgttacat ggccatggcg gtttccgcaa 180  
 ttggttttga aggtttcgag aaaaggttg aaatatcctt ttccagccgg gactttttgc 240  
 tgaccctgaa ggaaggggtc taagggctct tacaaaatcc cagttgggtg agatctaaca 300  
 ccagcn 306

<210> 2582  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 2582



agtgcacgc acgcgtacgt aagctcgga ttcggctcga ngccttcaca gcattcggtg 60  
 cggcatttta atcgatttat ccaagcagga ctgaatgaac taatggagtc taaaggtggg 120  
 aaaaagaagt ctagtagtag tagtagtaaa tcattgttct acgaagctcc cctcggtaca 180  
 gcattgaaga cgtttagacca aacgggtggaa tcaagaanct tcagatctgc tgcttactcc 240  
 aactgcgctc gcaaaccctc ctgatatcca tccggttgat agttncattg catgcnacca 300

<210> 2583  
 <211> 292  
 <212> nucleic acid  
 <213> Glycine max

<400> 2583

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 tcctctcttt gcgatttgtt ttgatctgc ttctactgc ganagtatcn ctactgttac 180  
 atggccatgg cggtnnccgc aattggtttt gaaggtttcg agaaaagggtt ggaaatatcc 240  
 ttttnccagc cgggactttt tgctgaccct gaaggaaggg gtctaagggc tc 292

<210> 2584  
 <211> 153  
 <212> nucleic acid  
 <213> Glycine max

<400> 2584

catgcacgcg tacgtaagct cggaattcgg ctcgagctca gaaatgatca tgtcgactcc 60  
tatgtttctgt ctgagtcag tctctttgtt tatgcctaca agatcatcat caaaacctgt 120  
ggtactacaa agctactgct tgcaatccca ccg 153

<210> 2585

<211> 474

<212> nucleic acid

<213> Glycine max

<400> 2585

tnnaactnta cgcgcccagg taccggtcaa agaattcccg ggtcgacca cgcgtcngta 60  
cggctgcgag aagacgacag aagggtacgg ctgcgagaag acgacagaag gggacacgca 120  
actattttctg actacgtttt gctctacgcc tctccctctc tctcaaaaat cgttctcttc 180  
gatttttaggg ttttgttttg ctgctgcctc cgttcccccc ttctcataaa caacgcggtt 240  
tctcttctgc ttcgatatcta ttctttgctt ttggttttgg ttgattcaag gccttcacag 300  
cattcgttgc ggcatttttaa tcgatttatc caagcaggac tgaatgaact aatggagtct 360  
aaaggtggga aaaagaagtc tagtagtagt agtagtaaat cattgttcta cgaagctccc 420  
ctcgatata gcattgaaga cgtagacca aacggtggna tcaaagaaat tcaa 474

<210> 2586

<211> 80

<212> nucleic acid

<213> Glycine max

<400> 2586

cagccctatc cccatcgcaa cttttctgag gaagttgcta ttcttgatgg ctacttnggc 60  
aagcattgct nggnagnggg 80

<210> 2587

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2587

tcgcatgcan ncgtagnaat aagctcnana attcggctcg agggcntgga atgggtggtt 60

ctctngtnta ncaaaaacct gccaaagactt gtgactgtgg ttcacctaga tcaactctga 120  
 agtgctggaa agaggaagat gaagaagagt agttttctta agtgtcttta ttatgtcctg 180  
 cgaaaataag tccggttttc cagacagtga ttgtttttct ttgggtgtttt ttccctttta 240  
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 gaa 303

<210> 2588  
 <211> 267  
 <212> nucleic acid  
 <213> Glycine max  
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ctgcacgcgt acgtaagctc ggaattcggc tcgagggaaa caagtcgttt gagcagacct 60  
 gctttctgga tgtaagggga tactgtncgt gaagagagga gccacgaagg gcttggaatg 120  
 ggtgggttctg ttgtctacca aaaattcggg aagactantg actgtgggtca cctagatcaa 180  
 ctctaaagtg ctggaatgag gaagatgagg aagagtagtt tccttaagtg tctttattat 240  
 tctgtccttg tgaaaataag tctgggtt 267

<210> 2589  
 <211> 225  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2589

ttcaatttct aggggttttgc tatacgcttt gcctccgttc cccctttctc acaaaaacaa 60  
 cgccttttct cttctccttc gtatctatc tttgcntttn gtttttggtt gattgaacgg 120  
 cattcacagc gtaattgggtg ctgcattttn atcgatttat ctaancagga ctgantgacc 180  
 taatggagtc taaaggtggg aanagaagt ctagtagtag tagta 225

<210> 2590  
 <211> 469  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2590

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agcccacgcg tccgtacggc tgcgagaaga cgacagaagg ggacacgcaa ctattttctga 120  
ctacgttttg ctctacgcct ctccctctct ctcaaaaatc gttctctgnc gatttttaggg 180  
tttcgttttg ctgctgcctc cgttccccc ttctcataaa caacgcgttt tctctttctgc 240  
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ggcattttta tcgattttat caagcaggac tgaatgaact aatggagtct aaaggtggga 360  
aaaagaagtc tagtagtagt agtagtaa atcattgttcta cgaagctccc ctcgataca 420  
gcattnaaag aacttngac caaacggtg aatcaaggaa attcagatc 469

<210> 2591  
<211> 298  
<212> nucleic acid  
<213> Glycine max

<400> 2591

cacatgcacg cgtacgtaag ctcggaattc ggctcgaggt tttattagga ttttntcttc 60  
tttgttttca attaggtttt ttgtgtgctc tccttcaaac tccatctttc caaatcctct 120  
ctttgcgatt gtgttttgat ctgcttcta ctgcgatagt ttctctactg ttacatgcc 180  
tggcggtttc cgcaattggt ttggaaggtt tcgagaaaag gttggaaata tccttttcca 240  
gccgggactt ttgtctgacc ctgaaggaag gggcttaagg gctctacaaa acccagtt 298

<210> 2592  
<211> 275  
<212> nucleic acid  
<213> Glycine max

<400> 2592

anacactctc ttgcaatttc tagggttttg gtattgcttt gcctccgttc cccctttctc 60  
ncaaaaaanaa cgnnttttct cttctccttc gtatctattc tttgcntntc gctttnggnt 120  
gantnaaggc attcanagcg taattngttn ctgcattttn atcgatttct ctaancagg 180  
ctgantganc taatggagtc taaaggtggg anaaagaagt ntagtagtan tagtagtaaa 240  
tgcnttttnt acgnagctcc cctcggatac agcat 275

<210> 2593  
<211> 269  
<212> nucleic acid

<213> Glycine max

<400> 2593

cgcgnnccgtn ngctcgggaat tcggctcgag cgcaaaccat cctgatacac atccggattg 60  
atagtccggt gcattgcaacc atagttttat taggattttt tcttctttgt tttcaattag 120  
gtttttttgt tgctctcctt caaactccat ctttccaaat cctctctttg cgattgtgtt 180  
ttgatctgct tctactgcg atagtntctc tactgttaca tggccatggc ggtttccgca 240  
attggttttg aaggtttcga gaaaagggtt 269

<210> 2594

<211> 155

<212> nucleic acid

<213> Glycine max

<400> 2594

acgncgcacc nacgcgtacg taagctcgga attcggtctg aggtgggtact acaaagctac 60  
tgcttgcaat cccacccata ttgaaatttg ctgaaatgct ttccctcaat gtttagatctg 120  
tgaattacac caggggaagt ttcattcttc ccggn 155

<210> 2595

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 2595

naaagtcgca ngcacgcgta cgtaagctcg gaattcggct cgagttcaaa tcacacactc 60  
tcttcaattt ctagggtttt gctattgctt tgctccggtt ccccttttct cacaaaaaca 120  
acgccttttc tcttctcctt cgtatctatt ctttgcgttt ngntttnggt tgattgangg 180  
cattcacagc gtaattngtt gctgcatttt aatcgattta tctaaccagg actgaatgac 240  
ctaattggagt ctaaagggtg gaaaaagaag tctagtagta gtagtagtaa atcaattttt 300  
n 301

<210> 2596

<211> 311

<212> nucleic acid

<213> Glycine max



<400> 2596

gtgcgatgca cgcnaacgta agctcggaat tcggctcgag gattgccttc aaatcacaca 60

ctctcttcaa tttctaggggt ttgctattg ctttgctcc gttccccott tctcacaaaa 120

acaacgcctt ttctcttctc cttcgtatct attctttgcg nttggttttt ggttgattga 180

aggcattcac agcgtaattn gttgctgcat ttnatcgat ttatctaacc aggactgaat 240

gacctaatgg agtctaaagg tgggaaaaag aagtctagta gtagtagtag taaatcaatt 300

tattacgaag c 311

<210> 2597

<211> 314

<212> nucleic acid

<213> Glycine max

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tctcttcaat ttctaggggt ttgctattgc tttgcctccg ntcccccttt ctcacaaaaa 120

caacgccttt tctcttctcc ttcgtatcta ttctttncnt ttngntttng gttgattgaa 180

ggcattcaca gcgtaattng ttgctgcatt tnnatcgatt tatctaacca ggactgaatg 240

acctaattgga gtctaaaggt gggaaaaaga agtctagtag tagtagtagt aaatcaattt 300

tttacgaagc tccc 314

<210> 2598

<211> 304

<212> nucleic acid

<213> Glycine max

<400> 2598

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ctattgcttt gcctccgttc cccctttctc acaaaaacaa cgctttttct cttctccttc 120

gtatctattc tttgcntttt gtttttggtt gattganggc attcacagcg taattngttg 180

ctgcatttnn atcgatttat ctaaccagga ctgaatgacc taatggagtc taaagggtggg 240

aaaaagaagt ctagtagtag tagtagtaaa tcaatctttt acgaagctcc cctcggatac 300

agca 304

<210> 2599  
 <211> 238  
 <212> nucleic acid  
 <213> Glycine max

<400> 2599

ctttctgatt gccttcaa at cacacactct cttcaatttc tagggttttg ctattgcttt 60  
 gcctccgttc cccctttctc acaaaaacaa cgccttttct cttctccttc gtatctattc 120  
 tttgcgtttg gtttttggtt gattganggc attcacagcg taattngtng ctgcatttna 180  
 atcgatttat ctaaccagga ctgaatgacc taatggagtc taaagnnggg aaaaagaa 238

<210> 2600  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max

<400> 2600

gtacgtnagc tcggaattcg gctcgagnag aaattcagat ctgctgctta ctctaactgt 60  
 tctcgcaa ac catcctgata cacatccgga ttgatagttc gttgcatgca accatagttt 120  
 tattaggatt ttttcttctt tgttttcaat taggtttttt tgttgctctc cttcaaactc 180  
 catctttcca aatcctctct ttgcgattgt gttttgatct gcttctact gcgatagttt 240  
 ctctactgtt acatggccat ggcgggtttcc gcaa 274

<210> 2601  
 <211> 257  
 <212> nucleic acid  
 <213> Glycine max

<400> 2601

caaganattc agatctgctg ctactctaa ctgttctcgc aaaccatcct gatacacatc 60  
 cggattgata gttcgttgca tgcaaccata gttttattag gatttnttct tctttgtttt 120  
 caattagggt tttttgttgc tctccttcaa actccatctt tccaaatcct ctctttgcga 180  
 ttgtgttttg atctgcttcc tactgcgata gtttctctac tgttacatgg ccatggcggt 240  
 ttccgcaatt ggntntg 257

<210> 2602

<211> 259  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2602  
  
 acgttttgcg ctacgcctct cctctctctt caaaaatcgt tcncttcgat tttagggttt 60  
 tgtttingctg ctgcctccgt tcccccttc tcataaaca cgcgttttct cttctgcttc 120  
 gtatctattc tttgcttttg gttttggttg attcaaggcc ttcacagcat tcgttgcggc 180  
 attttaatcg atttatccaa gcaggactga atgaactaat ggagtctaaa ggtgggaaaa 240  
 agaagtctag tagtagtag 259

<210> 2603  
 <211> 246  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2603  
  
 cgcacctott tctgattgcc ttcaaatac acactctctt caattttctag ggttttgcta 60  
 ttgctttgcc tccgttcccc ctttctcaca aaaacaacgc cttttctctt ctccttogta 120  
 tctattcttt gcgtttggtt tttggttgat tganggcatt cacagcgtaa ttngttgctg 180  
 catttnaatc gatttatcta accaggactg aatgacctaa tggagtctaa aggtgggata 240  
 nagaag 246

<210> 2604  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2604  
  
 annctcatgc acgcgtacgt aagctcggaa ttcggctcga gacgttttgc tctacgcctc 60  
 tccctctctc tcaaaaatcg ttctcttcga ttttagggtt ttgttttgcg gctgcctccg 120  
 tccccctt ctataaaca acgcgttttc tcttctgctt cgtatctatt ctttgctttt 180  
 ggttttggtt gattcaaggc cttcacagca ttcgttgcgg cattttaatc gntttatcca 240  
 agcaggactg aatgaactaa tggagtctaa aggtgggaaa aagangtcta gtagtagtat 300  
 agtaaatcat 310

<210> 2605  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2605  
  
 acgtcgcang cacgcgtacg taagctcggg attcggctcg agctacgcct ctctctctctc 60  
 tcaaaaatcg ttctcttcga tttnagggtt ttgttttgcg gctgcctccg ttccccctt 120  
 nctcataaac aacgcgtttt ctcttctgct tcgtatctat tctttgcttt tggttttggt 180  
 tgattcaagg ccttcacagc attcgttgcg gcattttaat cgatttatcc aagcaggact 240  
 gaatgaacta atggagtcta aaggtgggaa aaagaagtct agtagtagta 290

<210> 2606  
 <211> 333  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2606  
  
 nanatgnacg cgtacgtaag ctcggaattc ggctcgagct ttctgattgc cttcaaatac 60  
 cacactctct tcaatttcta gggttttgct attgctttgc ctccgttccc cttttctcac 120  
 aaaaacaacg ctttttctct tctccttcgt atctattctt tgcgtttggt tttcggttga 180  
 ntgaaggcat tcacagcgta attngttgct gcatttnnat cgatttatct aagcaggact 240  
 gantgagcca atggngtcta aangtgggaa aaagaagtct agtngtagta gtagtaaata 300  
 aattttttac gaagctcccc tcggatacag cat 333

<210> 2607  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2607  
  
 gcacgcgtac gtaagctcgg aattcggcnt cgagngattg cttcaaatac acacactctc 60  
 ttcaatttct agggttttgc tattgctttg cctccgttcc ccctttctca caaaaacaac 120  
 gccctttctc ttctccttcg tatctattct ttgcgtttng ctttngggtg actganggca 180  
 ttcacagcgt aattngttgc tgcattttna tcgatttatc taancaggac tgantgacct 240

aatggagtcc aaaggtggga aaaagaagtc tagtagtagt agtagtaa at caatTTTTTTa 300  
cgaagcnc cc ctg 313

<210> 2608  
<211> 286  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2608

accangcacg cgtacgtaag ctcgganttc ggctcgagtc tacgcctctc cctctctctc 60  
aaaaatcggt ctcttcgatt ttaggggtttt gttttgctgc tgccctccgtt ccccccttct 120  
cataaacaac gcgtttttctc ttctgcttcg tatctattct ttgcttttgg ttttggttga 180  
ttcaaggcct tcacagcatt cgttgcgga ttttaatcga tttatccaag caggactgaa 240  
tgaactaatg gagtctaaag gtgggaaaaa naagtctagt agtagt 286

<210> 2609  
<211> 311  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2609

ncgtcgang cacgcgtacg tnagctcggn attcggtcgc agnnttgctc tacgctctc 60  
cctctctctc aaaaatcggt ctcttcgatt ttaggggtntt gttttgctgc tgccctccgtt 120  
cccccttct cataaacaac gcgtttttctc ttctgcttcg tnttattctn gcgttttgg 180  
gatnggttga ntnaaggcct ncanagcatt cgntgcgga ttctaatacga nttatccaan 240  
caggntgaa tnaactantg gagtctaaag gnnggacaaa gaagtctagt ngtngtagna 300  
gtaaancatt g 311

<210> 2610  
<211> 306  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2610

ngtcnnangc acgcntacgt nagctcgga ttcggtcga gntctgacta cgttttgctc 60  
tacgcctctc cctctctctc aaaaatcggt ctcttacgat tttaggggtt ngttttgctg 120

ctgnectcgn tcccccttg ctcataaaca acgcgttttc nontctgctt cgtatctatt 180  
 ctttgcntnn ggntttgggt gattcaaggc cttcacagca ttogtngcgg catttnaatc 240  
 gatttatcca agcangactg aatgnactaa tggagtctaa aggtgggaaa aagaagtcta 300  
 gtagtg 306

<210> 2611  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2611

gtcgcangca gcgtacgtaa gctcggaatt cggtcgcagc gcctctccct ctctctcana 60  
 aatcgttctc ttcgatttta gggtnntgtt ttgctgctgc ctccgttccc cctttnctca 120  
 taaacaacgc gttttctctt ctgcttcgta tctattctac gcttttggtt ttgggttgatt 180  
 caaggccttc acagcattcg ttgnggcatt tnaatcgatt tatccaagca ggactgaatg 240  
 aactnatgga gnctaaaggn gggaaaaagc agtnntagtag cngtagcagt nacncatgtn 300  
 ntcagnagn cccngc 316

<210> 2612  
 <211> 329  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2612

acttcgcatg cacgcgtacg taagctcgga attcggctcg agctaacacc agctgcttgc 60  
 accattgttt cttcgctcan aancnntnaa tgtcgantcc tatgttctat ntgngtccag 120  
 cctgctttgt ttatgtcnta naagatcatc atcgcaacct gtggtntctac tangctattg 180  
 cttgnaatcc cacccatatt gnngttcgct ganatgcttt ccctgaatgt taagtctgtg 240  
 aattacacca agggaagttt cattttnnnc atgctcagnc atntccccgt cgcaactttt 300  
 ctgaggaagn tgctattctt ggatggcta 329

<210> 2613  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max

<400> 2613

agttctcatgc acgcntacgt nnagctcgga attcggctcg agtgcaacca tagttttatt 60

aggattttttt cttcttttgtt ttcaattagg tttttttgtt gctctccttc aaactccatc 120

tttccaaatc ctctctttgc gattgtgttt tgatctgctt cctactgcga tagttttctct 180

actgttacat ggccatggcg gtttccgcaa ttggttttga angtttcgag aaaaggttgg 240

aaatatcctt tttccagccg ggactttttg ctga 274

<210> 2614

<211> 275

<212> nucleic acid

<213> Glycine max

<400> 2614

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acacactctc ttcaatttct aggnnttttgc tattgctttg cctccgttcc ccttttctca 120

caaaaacaac gccttttctc ttctccttcg tatctattct ttgcgttttg tttttggttg 180

attgaaggca ttcacagcgt aatngtttgc tgcattttta tcgattttat taaccaggac 240

tgaatgacct aatggagtct aaagntggga aaaag 275

<210> 2615

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2615

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tcttcaattt ctagggtttt gctattgctt tgccctcggt ccccttttct cacaaaaaca 120

acgccttttc tcttctcctt cgtatctatt ctttgcgctt ggcttaggnt gactganggc 180

attcacagcg taattcggtg ctgcatttta atcgatttat ctaaccaggc ctgantganc 240

tantgggctc taaaggtggg aaaaagaagt ctaagtagta gtagagtaaa tcaantnncc 300

nc 302

<210> 2616

<211> 294

<212> nucleic acid

<213> Glycine max  
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 ttgcttttng ttttggttga ttcaaggcct tcacagcatt cgttgcgga ttttaatcga 180  
 tttatccaag caggactgaa tgaactaatg gagtctaaag gtgggaaaaa gaagtctagt 240  
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<210> 2617  
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 ctttctcaca aaaacaacgc ctttntctt ctcttcgta tctattcttt gognntngnt 180  
 ntnggttgat tganggcatt cacagcgtaa ttngttgctg cattttaatc gatttatcta 240  
 accaggactg aatgacctaa tggagtctaa anntggggaa aangaagtct agtagtagta 300  
 gtagtaaatac aatttntacg 320

<210> 2618  
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 <212> nucleic acid  
 <213> Glycine max  
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 ggccttcaca gcattcgttg cggcatttta atcgatttat ccaagcagga ctgaatgaac 180  
 taatggagtc taaaggtggg aaaaagaagt ctagtagtag tagtagtaaa tcattgttct 240  
 acgaagctcc cctcggatac 260

<210> 2619



<211> 285  
 <212> nucleic acid  
 <213> Glycine max  
  
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 cnatctcaaa aatcgtgctc ttcgatttta gggttttggn ttgctgctnc ctccgttcnc 120  
 cccttctcat aaacaacgnc gttttctctt ctgnttcgta tctattcttt gcttttggtt 180  
 ttggttgatt caaggccttc acagcattcg gtgcggcatt ttaatcgatt tatccaagca 240  
 ggactgaatg aactaatgga gtctaaaggt gggaaaaaga agtct 285

<210> 2620  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2620  
  
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 gctgcctcgc ttccccctt ctcataaaca acgcgttttc tcttctgctt cgtatctatt 180  
 ctttgctttt ggttttggtt gattcaaggc cttcacagca ttcgttgagg cattttaatc 240  
 gatttatcca agcaggactg aatgaactaa tggantcnaa nggtgggaaa aagaagtcta 300  
 gtag 304

<210> 2621  
 <211> 405  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2621  
  
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 tgtgaggtag actcgtggaa gcttcatttt ccctggggca cagtcttttc ctcaccgcag 120  
 tttttccgag gaggtttctg ttcttgacag ctatttcagc aaccttggtt ctggtagcaa 180  
 agcatatggt atgggtgacc cttcaaagtc acagatttgg cacatctact ctgcaagtgc 240  
 acagacaaaa ggatcatctg aagctgtcta tggcctagag atgtgcatga ccggtttaga 300

caaggaaagt gcttctgtgt ttttcaagga gaatacatct tcagcagctt caatgaccga 360  
aaattctgga attangaaga ttcttccaca gtctgatata tctga 405

<210> 2622  
<211> 299  
<212> nucleic acid  
<213> Glycine max  
<400> 2622

gtcgcatgca cgcgtacgta agctcggaat tnggctcgag ctcgagccgc cctgtcattc 60  
tcaagttggc tgatgctctt gacatagctg tgaaatctgt gaggtacact cgtggaagct 120  
tcattttccc tggggcacag tcttttcttc accgcagttt ttccgaggag gtttctgttc 180  
ttgacagcta ttccagcaac cttggttctg gtagcaaagc atatgttncg ggtgacctt 240  
caaagtcaca gatttggcac atctactctg caagtcaca gaccaaagga tcatctgaa 299

<210> 2623  
<211> 200  
<212> nucleic acid  
<213> Glycine max  
<400> 2623

ngtacgtaag ctcggaattn ggctcgagnn tcaagttggg ntgatgctct tgacatagct 60  
gtgaaatctg tgaggtacac tcgtggaagc ttcatTTTTCC ctggggcaca gtcttttctc 120  
caccgcagtt ttccgagga ggtttctgtt cttgacagct atttcagcaa ccttggttnt 180  
ggtagcaaag catatgttat 200

<210> 2624  
<211> 328  
<212> nucleic acid  
<213> Glycine max  
<400> 2624

gtnncatgca cgcgtacgta agctcggaat tcggctcgag ctgagaatgg gatggtgagt 60  
agtgaggggtg aaaccagttt cacttggtgc atgaagtttg gtggctcctc tgtggcttct 120  
gctgatagga tgaaagaggt ggctaccctt atattgagtt ttcccagga gaggcctatt 180  
gttgttctct ctgctatggg aaaaacaaca aacaagcttt tgctggctgg agagaaagct 240

gtgagttgtg gtgttatcaa tgtatcaagt attgaggagc tttgctttat aaaagacctg 300  
catctaagga ctgtggatca gcttggtg 328

<210> 2625  
<211> 254  
<212> nucleic acid  
<213> Glycine max  
<400> 2625

caaatgcgga catttttgaa gcaacttatc cggcagtcgc caagagatta catggtgatt 60  
ggctctctga tcttgcaatt gcaattgtta caggcttcct tggaaaggcc cagaaatcat 120  
gtgcagtgac aacactgggt agagggggca gtgatttgac agctacagca attgggaaag 180  
cactaggggt acctgagatc caggtatgga aggatgttga tgggtgccta acctgtgatc 240  
caaatatata ccca 254

<210> 2626  
<211> 297  
<212> nucleic acid  
<213> Glycine max  
<400> 2626

gagtcgcang cacgcgtacg taagctcgga attcggctcg agcaatattc gaagagttag 60  
gtatatcagt tgatgttgta gctacaagtg aagttagtat ttccttgaca ttggatccat 120  
caaagctatg gagcagagaa ctaattcaac aggaacttga ctatgttgtc gaagaactgg 180  
aaaaaattgc agtagtaaat ctcttaaaga ccagatccat aatctctctc attggaaatg 240  
ttcagagatc atcactaata ttggagaagg cctttcatgt tcttcgaact cttgggg 297

<210> 2627  
<211> 299  
<212> nucleic acid  
<213> Glycine max  
<400> 2627

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agtattgagg agctttgctt tataaaagat ctgcatctaa ggnctgtgga tcagcttggt 120  
gtggacggat ctgttattgc aaagcatcta gaagaattgg agcaacttct gaaggggata 180

gctatgatga aagaattgac taaaaggact caggactatt tagtctcctt ggagaatgca 240

tgctcgactag gatcttgctg gcatatctta acaaaatagg tgtcaaggcc gccatacga 299

<210> 2628

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 2628

tncacgcgta cgtaagctcg gaattcggct cgagcagata tgatgtatga ccttatccac 60

aaggctcaat caanagatga gtcttataca gctgcattaa atgctgtttt ggagaagcac 120

agtgcgaactg cacatgacat acttgancgg agataatctt gctactttct tgtctaaatt 180

gcatcatgat attagtaacc ttaaggcgat gcttcgtgca atatacatag ctggtcatgc 240

aacagagtcc ttacagattt tgttggtggga catggagaat aggtct 286

<210> 2629

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 2629

gtcgcgtacg tnagctcgga attcggctcg agcaggcctc aacctgaaa gaagaattca 60

acattgattt gcgtgtaatg ggcatacttg gttcanagtc aatgcttctt agtgatgtgg 120

gcattgactt agctagatgg agagaacttc gagaggaaag aggagaagtg gctaattgtgg 180

aaaaatttgt tcaacatgta catggaaatc attttatacc aaacacagca ttagtggact 240

gcacagctga ctctgccatt gctggctatt actatgactg gttgcgcaa 289

<210> 2630

<211> 168

<212> nucleic acid

<213> Glycine max

<400> 2630

angcacgcgt acgtnagctc ggaattcggc tcgagncaa atggatggat acaagggatg 60

tccttatcgt aaatcctact ggttctaatac aagttgatcc tgactatttg gaatctgagc 120

aaagacttga aaaatggtac tctttgaatc catgtaaggt aatcattg 168

<210> 2631  
 <211> 207  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2631  
  
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 ggctctgcat ccnngtcta tgagacctgc tanagaaagt gatattcctg ttaggggttaa 120  
 aaattcctac aaccctaaag ctccaggtac tctcnttgc tnnngacgngg gatatggnc 180  
 gggctttttt acctnccttt tttggag 207

<210> 2632  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2632  
  
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 aacgagttac acgtgtgtca tgaagtttgg cggctcctct gttgccaatg cagaaaggat 120  
 gagagaggtt gccaacctta ttctgagctt cccggaagag aggcctataa ttgtntcttc 180  
 tgccatggga aagacaacta acatgctgtt gctggctgga gaaaaagctg taagctgtgg 240  
 nataactang gnngatagtt tgacgnanng gttttnaaaa attggnatcc ggg 293

<210> 2633  
 <211> 270  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2633  
  
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 tctgttcaca aatttgggtg aacctgtgtg ggaacctctc aaagaataaa aatgtttgct 120  
 gacataattc ttaaggatga ttcggggaga aaattgggtg ttgtctctgc aatgtcaaag 180  
 gtgaaaaata tgatgtatga ccttatccac aaggctcaat cagcgatga gtcttatata 240  
 gctgcattgg attctgttta ggagaagcac 270

<210> 2634

<211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2634  
  
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 agtgacaaca ctgggaagag ggggcagtga tttgacggct acaacaattg ggaaagcact 120  
 agggttgcct ganatccagg tatggaagga tgttgatggt gtcctaacct gtgatccaaa 180  
 tatataccca aaagcagaac ctgttcctta tttgacattt gatgaggctg cagaactagc 240  
 ttacttttggg gctcaggttc tacatccaca gtctatgaga cctgccagag aaagtgatat 300  
 tcctgttagg gtt 313

<210> 2635  
 <211> 322  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2635  
  
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 tagtntcctt tggagaatgc atgtnnacaa ggatctttgc nggnacatnc gnaanaaaat 120  
 aggtgtcaag gctcgccaat atgatgcatt tganattggt tttataanca actgacgact 180  
 tcacanatgc ggacattttg gnngccactt atccagctgt tgcanagagn ttgcatggtn 240  
 antagctctc cgctcctgca attgcaattg cnacggctcc ttggaaaggc ccggaactca 300  
 ngtgcantnc caacactggg ac 322

<210> 2636  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2636  
  
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 tccttgacat tggatccatc aaagctatgg agcagagagc taattcagca ggaacttgac 120  
 catgtttgtag aagaactcga gaaaatcgct gtgggtgaatc tcctgcagaa tagatccatc 180  
 atctctctca ttggaaatgt tcagagatca tcactaatat tggagaagggt tctgtattat 240

gctcccttta tttaaattag ttcatatcct aactttcact tatgtataaa gatttactga 300  
 atttattaca 310

<210> 2637  
 <211> 438  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2637

ccacgcgtcc gtcccaaata tggaagaata tcttggaag caggtcttag aggctgtttc 60  
 attgattgaa gttagaagga acttcattga agcattatcc agtccgtttg gaaggccagt 120  
 tgaagctgat gctgtctttt gcaggaagga aacttttctt gctgcatctg gtgttttcac 180  
 gttcctgaag cagaagaaat accgtgtgac agaaatgttg ggatccaatt gcaagatatg 240  
 ggacttgagt tccacattgg aagtaatgag attctaattg gggttttata agacottggg 300  
 ctctctgact acaatgatta gcttttgtgg tatgattcac tcaaagttct tatcaattgg 360  
 tattagagct tttcattatg tctntcaact ncaaaacanc nacntgggtg aataccttca 420  
 aatttgnact gttnaagt 438

<210> 2638  
 <211> 329  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2638

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 attcaggacc acttgatcag tatttgaagt taagagctct tcaaaggcaa tcctatacac 120  
 attacttcta tgaagcaact gtcggagctg gtcttccaat tgtagcact ttacgtggcc 180  
 tccttgaaac tggagacaaa atattacaaa tcgaaggcat ctttagtggg actttgagtt 240  
 acatatttaa taactttaaa gatggccggg ctttagtga ggtagtttct gaagcaaagg 300  
 aagcaggtta tactgagcca gatccaaga 329

<210> 2639  
 <211> 256  
 <212> nucleic acid  
 <213> Glycine max

<400> 2639

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actgttggag ctgggtcttcc aattgttagc actttacgtg gcctccttga aactggagac 120  
aaaatactgc aaatcgaagg catcttttagt gggactttga gttacatatt taataacttc 180  
aaagacggcc gggcttttag tgaggtagtt tctgaagcaa aggaagcagg ttatactgag 240  
ccagatccaa gagatg 256

<210> 2640

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 2640

gtcgnnntnn gggtagctga gctcggnatt cggctcgagg ggcagtatta acaagcattg 60  
ttttgaaacg taatgtgacc atgttggata tagcaagcac tcgcatgctt ggtcngtatg 120  
gtttccttgc taaggngttt tcaatctttg aagagttagg catatcagtt gatgtttag 180  
ctacaagtga agtcagtgtt tccttgacac tggnnccatc aaagctatgg agcagagagc 240  
taattcagca ggcaagtga cttgaccatg ttgtagaaga actcg 285

<210> 2641

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 2641

cgcngcnttg tacgtnagct cggnatctcg ctcgagggca gtattaacaa gcattgtttt 60  
gaaacgtaat gtgaccatgt tggatatagc aagcactcgc atgcttggtc agtatggttt 120  
ccttgctaag gtgttttcaa tctttgaaga gttaggcata tcagttgatg ttgtagctac 180  
aagtgaagtc agtgtttctt tgacactgga tccatcaaag ctatggagca gagagcta 240  
tcagcaggca agtgaactga ccatgttgta gaagaactcg ag 282

<210> 2642

<211> 527

<212> nucleic acid

<213> Glycine max



<400> 2642  
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 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nncgagcaat aattgttctc tctgccatgg 120  
 gaaagacaac taacatgctt ttgctggctg gagaaaaagc tgtaagctgt ggagtaacta 180  
 atgctgatag tattgatgag ctcaacatag taaaagatct tcatctcagg actgtggaac 240  
 agcttggagt ggacagaaat gttattgaga agcatctaga agaattggag caacttctaa 300  
 aggggatagc tatgatgaaa gagttgactc cacggactca agactattta gtttcatttg 360  
 gagagtgcac gtccactagg atatttgctg catatcttaa tacattagga gttaaggccc 420  
 gccaatatga tgcatttgag atgggtatta taacaactga tgacttcaca aatgctgaca 480  
 ttttggaagc aacatatact gctgttgcaa aaaggttcat aantgat 527

<210> 2643  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max

<400> 2643  
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 gcttcccgga agagaggcct ataattgttc tctctgccat gggaaagaca actaacatgc 120  
 tgtngctggc tggagaaaaa gctgtaagct gtggagtaac tatggctgat agtattgacg 180  
 agctcagcat tataaaagat ctgcatctca ggactgtgga agagcttgga gtggacagaa 240  
 atgctattga gaagcatcta gaagaattgg agcaactttt aaaggggata g 291

<210> 2644  
 <211> 265  
 <212> nucleic acid  
 <213> Glycine max

<400> 2644  
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 aaccttattc tgagcttccc ggaagagagg ccaataattg ttctctctgc catgggaaag 120  
 acaactaaca tgcttttgct ggctggagaa aaagctgtaa gctgtggagt aactaatgct 180  
 gatagtattg atgagctcaa cataatanaa gatcttcatc tcaggactgt ggaacagctt 240

ggagtggaca gaaatgttat tgaga

265

<210> 2645  
<211> 307  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2645

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cccttttcaa tgcattggct aaggccaata taaatgtccg tgctatagcg caaggttggt 120  
ctgagtacaa tattactggt gttgttaagc gagaggattg tataaaggct ttacgagctg 180  
tccattccan attttatctc tcaagaacca ccatagcaat gggcattatt ggacctggat 240  
taattgggag cacactactt gaccagctaa gggatcaggc ctcaaccctg aaagaagaat 300  
tcaacat 307

<210> 2646  
<211> 327  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2646

nngnngcann cagcggtacg tnagctcgga attcggctcg anntccgtgc tatagcgcaa 60  
ggntgttctg agtacaatat tactgttggt gttaancgag aggnttgat aaaggcttta 120  
cgagctgtcc attccagatt tnatctctca agaaccacca tagcaatggg cattattgga 180  
cctggattaa ttgggagcac actacttgac cagctanagg atcaggcctc aacctgaaa 240  
gaagaattca ncattgattt gcgtgtaatg ggcatacttg gttcaaagtc aatgctotta 300  
gtgatgttgg cattgacttn ncctagn 327

<210> 2647  
<211> 317  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2647

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aagaattgga gcaacttctg aaggggatag ctatgatgaa agaattgact aaaaggactc 120

aggactatTT agtctcctTT ggagaatgca tgtcgacaag gatctttgct gcataTctta 180  
 ataaaatagg tgtcaaggct cgccaatatn atgcatttga gattggTTTT ataacaactg 240  
 acgacttcac aaatgCGgac atTTtggaag cnaCTtatnc agctgttgca aagagattgc 300  
 atggtgattg gctctcc 317

<210> 2648  
 <211> 334  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2648

tcgcangcac gcgtacgtaa gctcgggaaT tcggctcgag ctcaTTtcca gagcatctag 60  
 aagaattgga gcaacttctg aaggggatag ctatgatgaa agaattgact aaaaggactc 120  
 aggactatTT agtctcctTT ggagaatgca tgtcgacaag gatctttgct gcataTctta 180  
 ataaaatagg tgtcaaggct cgccaatatg atgcatttga gattggTTTT atagcaactg 240  
 acgacttcac aaatgCGgnc atTTtggaag caacttatcc agctgttgca aagagattgc 300  
 atggtgattg gctctccgat cctgcaattg caat 334

<210> 2649  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2649

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 tccatcgccg tcgtgggCGt catcggcgCC gncggccagg agttcctact cgtcctctc 120  
 cgaccgCGac ttccctacc gctcctcat atgctggctt ccaagcGctc cgtgggccgc 180  
 cgcacacct tcgaggacag ggactacGtc gtccaggagc tcacgCCgga gagcttcGac 240  
 ggtgtcgcat cgcGctcttc agcGccggcg gctccatcag caagca 286

<210> 2650  
 <211> 280  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2650

gtnnangcac gcgtacgtaa gctcgggaatt cggctcgagg acgctccggt gggcgccgca 60  
tcacctttna gnnccagggga ctacgtcgtg nccgagctga cggcgggagan cttcgatggc 120  
gtcganacgc cgatctncan cgcgcncggc nccattagc aagtactttn gccccatcnn 180  
cgtnatcgg ggaacgggtgn tcgncgacan cagatccgcn tntcggatgg acnanaatgt 240  
cccattggta atncccgaat caaaccgtn nccatgcaaa 280

<210> 2651  
<211> 323  
<212> nucleic acid  
<213> Glycine max

<400> 2651

ncacgcgtnc gtanacgtnn gnattcngct cgannngcgn nnnngctgggg cgccgcatca 60  
cctttgagtn acaggggacta cgtcgtggag gagctgacgg cggagagctt cgatggcgtc 120  
gacatcggcg tottcagcgc cggcgggctcc attagcaagt acttcggccc catcgccgtc 180  
gatcggggaa cgggtggctgt cgacaacagc tccgcgtttc ggatggacga gaatgtccca 240  
ttggtaatc ccgaagtga cccggaagca atgcaaaa tcaaagccgg aatggaaagg 300  
gcgcatcat gctaacccta att 323

<210> 2652  
<211> 402  
<212> nucleic acid  
<213> Glycine max

<400> 2652

gggaataaaa cctgctttct atttttctca acctaaaatc ccatccacca acottatgat 60  
gatgttttca ctctctgttt cgcgccacaa ccacctcttc tcgggccctc tcccggcccg 120  
gcccagccc aagcccagct tttcctcttc caggatccga atgtccctcc aagaaaacgg 180  
cccctccatc gccgtcgtgg gcgtcaccgg cgcgcgcggc caggagtctc tctcgtcct 240  
ctcgcaccgc gacttccctt acagctccat caaaatgctc gcgtccaagc gctccgctga 300  
gcgcgcgcatc acctttgagg acaggggacta cgtcgtggag gagctgacgg cggagaactt 360  
cgatggcgtc gacatcgcg ctttcancgc cggcgggctc aa 402

<210> 2653

<211> 482  
 <212> nucleic acid  
 <213> Glycine max

<400> 2653

gggnngggnn nnnnttaact ttccagggcc cggtcaggaa aacccgggtc gacccacgcg 60  
 tngtacggc tgcgagaaga cgacagaagg gggggacatg gaattggaag acaagttgga 120  
 gtgtggtggt ctgttttaaa atccaacact taatctctct cttcgagcc taaaatccca 180  
 atggcttcac tctctgtttt gcgccacaac cacctcttct cgggccccct ccggccccgc 240  
 cccaagccca cctcctctc ctctccagg atccgaatgt ccctccgga gaacggcccc 300  
 tccatcgccg tcgtggggt caccggcgcc gtgggagcagg agttcctctc cgtcctctcc 360  
 gacggcgact tcccctaccg ctccattcat atgtgtggtt ccaagcgctc cgtgtggcgc 420  
 cgcattacct tcgagacaa ggactaagtc gtccaagaac tcaggccggg anaacttcaa 480  
 cg 482

<210> 2654  
 <211> 327  
 <212> nucleic acid  
 <213> Glycine max

<400> 2654

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 atggcttcac tctctgtttt gcgccacaac cacctcttct cgggccccct ccggccccgc 120  
 cccaagccca cctcctctc ctctccagg atccgaatgt ccctccgga gaacggcccc 180  
 tccatcgccg tcgtggggt caccggcgcc gtgggagcagg agttcctctc cgtcctctcc 240  
 gacggcgact tcccctaccg ctccattcat atgtgtggtt ccaagcgctc cgtgtggcgc 300  
 cgcattacct tcgaggacag ggactac 327

<210> 2655  
 <211> 312  
 <212> nucleic acid  
 <213> Glycine max

<400> 2655

gtngcccnca cgcgtacgta agctcggaat tcggctcgag gagctttgcc acaagaaggg 60

gactttgctc tgtattgatg gtacatttgc aacaccattg aaccagaagg cccttgcctt 120  
 tggcgctgat ctgattctgc actccttaac aaaatacatg ggtggacatc atgatgtaag 180  
 ttggcatggt caagatttag ttgatggaa aaacaacaca aatatgttac attttcaggg 240  
 ttaagagtta agcgtcaagg agcattttcc tctttcaggg ccttgggtgg tgcataagtg 300  
 gttcaactaa gg 312

<210> 2656  
 <211> 542  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2656

gnnnangtnn ntntcnaana ngntaanctt tnnaaacctt ccatttnggt aanncnccggg 60  
 gccacgcgtc cgcncacgcg tncgcccacg cgtccggaga gaaagagagg gagtaagtgg 120  
 gtinnaggang gaaaactaaa gaaacaaacc taacacaaca caaatccttg aaacgacgac 180  
 ggaaatggcc gtttcgagct cgcacatgcg tttcaccttt gagtgccgct ccgatcccga 240  
 tttctcgcgc cccccgcgt ccttcgacaa cctccgccgc cgaaaacttc gctcctccgc 300  
 aggatccggc gcggcggttt acggcatctc ctccctcatc ctccgcttcc tcccaacttc 360  
 cagcgccagc taagcaccaa ggcgcgccgc aactgcagca acatcggcgt cgcgcaaata 420  
 gtccgcgctt cgtggtcgaa caacagcnac aactnttcgg ccgncggggc ttncgcgcgc 480  
 accgggggnc cgggcacgga cccgggtacg ggcttttccc gtngtcgtaa cgncaacaa 540  
 gg 542

<210> 2657  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2657

cgtcgcangc acgcgtacgt ncagctcgga attcggctcg agcggatctt gacatgggtt 60  
 atgctgccat agtagaaggg aaaacaaaag tgctttactt cgaatctgtt tccaaccca 120  
 cccttacggt tgcgaacata cctgaactgt gccacatggc acaccggaag ggagtgcg 180  
 tgggtggtgga caacacgttc gcgcccattg tgctttcgcc agcgcgtctt ggtgctgatg 240

ttgtcgttca cagtatctcc aagttcatca gcggtggggc cgatatcatt gcaggagcgg 300  
tgtgcg 306

<210> 2658  
<211> 307  
<212> nucleic acid  
<213> Glycine max  
<400> 2658

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gacggatctt gacatgggtg 60  
atgctgccat agtagaaggg aaaacaaaag tgctttactt cgaatctgtt tccaacccca 120  
cccttacggg tgcgaaacata cctgaactgt gccacatggc acaccggaag ggagtgaagg 180  
tggtgggtgga caacacgttc gcgcccattg tgctttcgcc agcgcgtctt ggtgctgatg 240  
ttgtcgttca cagtatctcc aagttcatca gcggtggggc cgatatcatt gcaggagcgg 300  
tgtgcg 307

<210> 2659  
<211> 515  
<212> nucleic acid  
<213> Glycine max  
<400> 2659

gnnnnngnaga gtttgnttgg ggggnaggga gnnanatttt nagaccacta tgacgtcgca 60  
tgcacgcgta cgtaagctcg ganttcggct cgagagacag aggntagaga angagaggga 120  
gtaagtgggt aaaggaaaga naactaaaga aacaaaccta acacaacaca aatccttgaa 180  
acgacgacgg aaatggccgt ttcgagctcg cacatgcgtt tcacctttga gtgccgntcc 240  
gatcccgatt tctcgcccc cccgccgtcc ttcgacaacc tccgccgcg caacttccgc 300  
tctccgcag gatcccgggc cggcggttca cggcatctcc tctcctcct ncgcttccct 360  
cccaacttcc agcgccagct aagcaccaag gcgcgccgca actgcagcaa catcggcgtc 420  
gcgcaaactc tcgccgcttc gtggtcgaac aacagcgaca actctccggc cggcggggct 480  
tcggcgccgg ccgcggcacc ggcacggacg ccgtt 515

<210> 2660  
<211> 258  
<212> nucleic acid

<213> Glycine max

<400> 2660

caaggacatc atgatgtcct tgggtggttgc ataagtgggt caattnnngt ggtttcgcaa 60  
attcggactt tgcaccatgt nttgggtggt aactttaacc cgaatgctgc atacctatc 120  
atcagaggca tgaaaacgct gcatctccgt gtacagcagc agaattcaac aggaatgagg 180  
atggccaaac ttttagagggc acatcccaag gtgaagcggg tctactatcc aggcttgccg 240  
agtcaccctg aacatgag 258

<210> 2661

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2661

ganattgtca tgcactctgc tacaaaatth attgctggac atagtgacat tatggctggt 60  
gtgcttgctg tgaaggggtga aaagtggga aaggaattgt atttcttgca aaatgcagag 120  
ggttcaggct tagcaccatt tgactgttgg ctttgtttgc gaggaatcaa gacaatggcc 180  
ctgcgaattg aaaaacaaca ggataatgca cagaagattg ctgagttcct tgctcccat 240  
cctcgagtga agaaagtga ttatgctggc ttgctg 277

<210> 2662

<211> 322

<212> nucleic acid

<213> Glycine max

<400> 2662

gcatgcacnc gtacgtaagc tcggaattcg gctcgagntt gcctcccatc ctcgagtga 60  
gaaagtgaat tatgctggct ngcctgggtca tcctggctgt gatttacact attctcaggc 120  
aaaggggtgca ggatctgtgc ttagcttctt gactggttca ttggaacttt caaagcatat 180  
tgttgaaact accaaatact tcagtataac cgtcagcttt gggagtgtga agtcccttat 240  
tagcatgcca tgctttatgt cacatgcaag cnngncngct ggcagttcgt gaggccagag 300  
tttaagaaga tctgtagtat at 322

<210> 2663



<211> 273  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2663  
  
 cgcangcacg cgtacgtaag ctcggaattc ggctcgagaa acaacaggat aatgcacaga 60  
 agattgotga gttccttgcc tcccatcctc gagtgaagaa agtgaattat gctggcttgc 120  
 ctggtcaccc tggctgtgat ttacactatt ctgaggcaaa ggggtgcagga tctgtgctta 180  
 gcttcttgac tggttcattg gaactttcaa agcatattgt tgaaaactacc aaataacttca 240  
 gtataaccgt cagctttggg agtgtgaagt ccc 273

<210> 2664  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2664  
  
 gcaagcggtac gtaagctcgg aattcggctc gaggttcagg cttagcacca tttgactggt 60  
 ggctttgttt gcgaggaatc aagacaatgg ccctgcgaat tgaaaaacaa caggataatg 120  
 cacagaagat tgctgagttc cttgcctccc atcctcgagt gaagaaagtg aattatgctg 180  
 gcttgccctgg tcctcctggg cgtgatttac actattctca ggcaaagggt gcaggatctg 240  
 tgcttagctt cttgactggt tcatggaact tcaaagcata ttgttgaaa 289

<210> 2665  
 <211> 499  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2665  
  
 aacttttacg cncangtg ccgggcaang gtnangagnt cccgggtcga cncacgcgtc 60  
 cggcnagaag acgacagaag gggacgggtg tgggtggacaa cacgttcgcg cccatgggtgc 120  
 tttgccagc gcgtcttggg gctgatgttg tcgttcacag tatctccaag ttcacagcg 180  
 gtggggccga tatcattgca ggagcgggtg gcggaccgcg aagactgggtg aacgcaatga 240  
 tggatctgca acaagggtca ctaatgctgc tgggtccaac aatgaatgcg aaagtggcat 300  
 tcgaactctc ggagagaata ccgcacctag ggctaagaat gaaggagcat agcaaccgcg 360

cactagagtt cgcaacgagg ctcaaaaggc taggaatgag ggtaatatatac ccgggcctgg 420  
 aggagcaccc acagcaccaa gcttctgaaa tcaatgcaca acaaggacta tggctacggc 480  
 gggctcatgt tcntngaca 499

<210> 2666  
 <211> 326  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2666

tgcgatgcac gcgtacgtaa gctcgggaatt cggtctgaga cgggaggtca agcccgcct 60  
 gaccaacatg gtctccgctg ctaagctcat tcgcacccag ctgcgcagcg ccaagtgagc 120  
 accttttttt gccttttttcg ttcccgagga gggcgctcgtc gatgcccaatt tgtntccaat 180  
 aaacaggggt cccccctgt gcccgccgtt ctgttgtgct ccgtctgtgg ttaggttact 240  
 agttttcttg atctcgcccc cacgcgggtac ctgttttata tctgtttgggt ggtttctgag 300  
 gcaagttgcc cgtgtattgt atcgta 326

<210> 2667  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2667

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gggacactac tcagattcac 60  
 actcacatgt gctactccaa cttcaatgac atcattcact caatcataga catggatgct 120  
 gatgtgatca ccattgagaa ctccagatca gatgagaagt tactttcgggt cttccgtgag 180  
 ggagtgaat atggtgccgg cattggtcct ggcgtttatg atattcactc acccaggatt 240  
 cctcccacag aagaaattgc tgacaggatc aacaagatgc ttgcagttct tgaaagcagc 300  
 attctctg 308

<210> 2668  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2668

nagtcgcacg cacgcttacg taagctcnga attcggetcn nnnngccant gancttgctg 60  
 cttctctcan taccttggca gggctcttgag ggcatgtgtg ggcaaagata agcttggttg 120  
 tgtccacctn ctctccctt cttcacactg gctgtggntc tagttaacga naccnagttg 180  
 gatgatgaga tcaagtcatg gctagctttg gctgccccaa aaattgttga ngttaacgca 240  
 tnggctaaag cngtgtcngg ccacaaggat gaggcctnct tctctgntan tgcngctgct 300  
 ctggcttcaa gan 313

<210> 2669  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max

<400> 2669

cgcangcacg cgtacgtaag ctcggaattc ggctcgaggg ggatgtactt gatcccannc 60  
 accagccatc tgctcgacgc caccgccanc ctcggtgccg tcccccccag gtacggcttg 120  
 accggcggcg agattggatt cgnacactac ttctccatgg ccagaggtaa tgctaccgtg 180  
 cntgctangg agatgaccaa gtggttcgac accaactacc actttattgt ccoctgaattg 240  
 ggccctgatg tgaattcacc tatgcttctc acaaggctgt tgatgaatac 290

<210> 2670  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max

<400> 2670

cncgcacgcg tacgtaagcn tcgcgaattc ggctncgagt tgctcccnca gnagcganga 60  
 agangccaca gagnnctagt ctccctanct ctacacccgc aaganaaana tggcatctca 120  
 catacgttgg atacccccgt atgggttccc aagagagnnt ncaanttcgc tctcgagtcn 180  
 ttctgggatn gcaanancag cgccgangct ttgcagangg tgtcntctga tctcagggca 240  
 nccatctggn ancagttggc tgatgntggg ntcaagtaca tccccngca 289

<210> 2671  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max

<400> 2671

nttgcgtnc a cgcgtacg nn agctcgg aat t cggctcgca gtgcacgtgt nattangnna 60  
aatgagaaan aaaaaaatgg tcatncaca tcnnggata cccccgtatg ggtgccaag 120  
agagagcnca agntcgcttc tcgagtcttn cngggatggc aagagcagcg cngaggattt 180  
cncagaaggt gtcttcngan ctcagggcat ccancnggaa gcagatggct gttgctggga 240  
tcaagtanan ccnancan acttcngcnc actatgatca ggntcacnac gncctgccac 300  
nct 303

<210> 2672

<211> 284

<212> nucleic acid

<213> Glycine max

<400> 2672

agngacgcgt actaagtcgg aattcggctc gagaagataa gcttggtgtg tccacctcct 60  
cctcccttct tcaactgct gtnganccca gttaacgaga ccaagttgga tgatgagatc 120  
aagtcattggg ctagcttttg nctggcccaa naaattgtn ngaagttaaa cggcattggg 180  
ctaaaaggca ttggtgctgg gaccancaa ggatggaggg cctttctttc tctgggtaaa 240  
tggttggtg gctactgggc tttccaaggg aaagttcttt cctn 284

<210> 2673

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2673

cntgcncgta nntnagctcg gaattcggct cgagctgagg ttgttggtgt tgccaaggnt 60  
gtaaatnatg gtcccgttc tgtgcgtgat ggctnctnaa nccaatgcca aatccatgca 120  
ggngentgcc acgtcgaanc gtaccantaa caaggcggtt aaggaccgtc aagccagcgt 180  
cactcctgag caacacngc gcaagtctca gttccctgaa cgttatgcc agcagaagaa 240  
gcacttgagc ctcttggttc cccaccacca ccattggttc cttccctcag accana 296

<210> 2674

<211> 269

<212> nucleic acid

<213> Glycine max

<400> 2674

gcgcntgcan tnanacgtng agntcggaat tcggctcgag cttccaatcc ttccaaccac 60  
cactattggn ncnttcngtc agantnnann antgaggagg gtacgccgtg agttcaangc 120  
taanaagntc tnnnnnggaa ntntatgnct aagtcaatta aggnnggaaat tcgcaaagtt 180  
gttgaacttc aagaagagct tgatattgat gttcttggtc atggagancc agagngaaat 240  
gntatgggtg agtacttcgg tgagcaatt 269

<210> 2675

<211> 216

<212> nucleic acid

<213> Glycine max

<400> 2675

nngtcganac ntgcgtacgt aancncggaa ttcggctcga gggcaagacc agcgccgnng 60  
attngcaggn ggtgtctnct aancncagga gcatccatch ggaagcagat ggcngatgct 120  
gggatcaant acatccccag caacactttg ctactcacna naaccaggtn ctcgacgccn 180  
ccgccaccct acggngccgn tgccacnnag gtangg 216

<210> 2676

<211> 263

<212> nucleic acid

<213> Glycine max

<400> 2676

anatgaccag aagacggtca ttggctttgg gcggctctgg ctcagagctt taccaagcgc 60  
ccaatgaagg gaatgcttac cggaccagtt gagaatggta acaggaccgg taagcattct 120  
caactgggtcc ttgtttagaa atgaccaacc tagatctgag accacctacc agattgcttt 180  
gtctatcaag gacgaagtgg aagacctga aaaggctggc atcactgtta tccaaattga 240  
tgaagctgct ttgagagagg gtc 263

<210> 2677

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 2677

cgcangcacg cgtacgtaag ctcggaattc ggctcgagtn gggngcgggg ggcgtgaagc 60  
caccgatcat nnatgggtgat gtgagccgcc caaagccaat ganngtcttc tngngatnnc 120  
tggntcagag ctttacnaag cgcccaatga agggaatgct taccggtcct gttaccannc 180  
tcaactgggn nnntgttaga aatgaccanc ctagatctga gaccacctag nagantgctt 240  
tgtctatcaa ggacgaatgg aaganncnhn ccaaggctgg catcantgtn a 291

<210> 2678

<211> 267

<212> nucleic acid

<213> Glycine max

<400> 2678

cctanggtgc cgntccacna nggnacngnt gncnccggcg gngagattng gtttgatacc 60  
tactttctcca tggccanang taatgctacc gtgccagcta tggagatgac caagtgggttc 120  
gacaccaact accactntat tgtccctgaa ttggggccctg atgtgaactt cacctatgct 180  
tctcacaagg ctgttgatga atacaaggag gccaaaggcg ttgnagtga taccgttccg 240  
gtcctcgttg gccctgttac atacctg 267

<210> 2679

<211> 252

<212> nucleic acid

<213> Glycine max

<400> 2679

cagaagccnc agaagaagcc acagagaact agtctnctac tcnccaccng caagnaccnn 60  
natggcatnn tcnctcggtt ggataccccc gtatgggtcc ncaannngag agtcaagtt 120  
gtctcgagtc tttctgggat ggcaagagnn agcgccgagg gatttgcaga aggtgtcttc 180  
tgatctccag ggcattccatc tggaagcaga tggatgatgct gggatcaagt acatccccag 240  
caaactttct nt 252

<210> 2680

<211> 324

<212> nucleic acid

<213> Glycine max

<400> 2680

gtcgcangca cgcgtacgtn agctcggnat tcggctcgag cacgagacca agttggatga 60  
tgagatcaag tcatggctag cttttgctgc ccaanaaatt gttgaagtta acgcattggc 120  
taaagcattg tctggccaca aggatgaggc cttcttctct ggtaatgctg ctgctctggc 180  
ttcaaggaag tcttctccaa agagttgacc aaacgagggc tgntccagaa agnctgctgc 240  
tagcaattga agggttccag atcatngccg gncaacaatt ntccatgcc aactggatnc 300  
tcaacaaaag aagnncaacc ttcc 324

<210> 2681

<211> 362

<212> nucleic acid

<213> Glycine max

<400> 2681

tnnaggtagt agctcggatt cggctcgagct ctccatggcc agaggtaatg ctaccgtgcc 60  
tgetatggag atgaccaagt ggtncgacac naactaccac tttattgtgc ccngnattgg 120  
gccctggatg nagaactttc acctatgnct ntcttcacaa gggctgtntg gatgcagata 180  
ncaagggaag gtccaagggc cgatttgagg agtcgggata nccaatntcc ncggnaactc 240  
cgtntggggc cnetgggtta ancatnactt tggtataggc tacctnncca aaggctcctg 300  
gcacaaangg ggcaaatttc gganggtaaa attcccattg gggggcttcc tggccttgc 360  
tc 362

<210> 2682

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2682

cgtegnange ncgcgtncgt cagctcggnn ttccgctcgn gctgnacctg tgcttgtggg 60  
acctgtttct tncctgttgc tgtcannacc agctaagggt gttgagangt cattttccct 120  
tctttcccta attgacaaga tcttctctgt ctacaggga gttgtggctg aactgaaggc 180  
agctggtgct acttgatcc agtttgatga acctaccctt gtgaaggatc tcaatgcccc 240  
ccagttacaa gcatttacc atgcctacgc agagtttagag tcaagtttnt ctggtttggn 300

tgttctgatt ganacatact t

321

<210> 2683  
<211> 315  
<212> nucleic acid  
<213> Glycine max

<400> 2683

gnngcacgcg tacgtaagct cggaattcng ctcgagctgn cntcaacttg ntccctcaga 60

agcgnntann aatccacana gaactatgtc tcnctactnc ncacccgcaa gaaaaaaatg 120

gcntntcaac atcgttggat acccccgtat nggncccaat agaganctna attcgctctc 180

gagtctatct gggatgncaa nagtacgccg aggatgttga naangngtct actgatctca 240

gggcattcnt ctgganncag atggntnatg ctgggatcag tacatccnca gcaacatttc 300

tctnactntg accan 315

<210> 2684  
<211> 174  
<212> nucleic acid  
<213> Glycine max

<400> 2684

gctgcaggca cgcgtacgtn agctcggaat tcggctccgn gcttcanctt gctccctcag 60

aagcgnagaa gaagcccana gagaanagnc tcctantctc acccgcaagn nnaanatggn 120

atctcacatc gttggntacn cccgtatggg tcccaagaga gagctnaagt tgnt 174

<210> 2685  
<211> 303  
<212> nucleic acid  
<213> Glycine max

<400> 2685

aacngaaagn cgcangcacg cgtacgtaag ctcggaattc ggctcgagct tcaggatctc 60

cgaggaagag tatgtaaagt caattaagga ggaaattcgc aaagttgttg agcttcaaga 120

agagcttgat attgatgttc ttgttcattg agaaccagag gtccgctctc atttcataac 180

atgactaaat attagtcttt tgaattgaag atagcttctt tctttctgaa gagcaactac 240

tctttgcata ttttctttcc tattgttaga ttgcngattc caaaattgca gcacctccaa 300



ctn

303

<210> 2686  
 <211> 103  
 <212> nucleic acid  
 <213> Glycine max

<400> 2686

gntttntttt ttttttgggt tgttttcnnt tggttttttt tgtttttnnt gnttnttggt 60

tnnttggttt tttttgtttt ttntgtttt ttttttttt nnt 103

<210> 2687  
 <211> 523  
 <212> nucleic acid  
 <213> Glycine max

<400> 2687

gnnngnaggt tttgannggg ggggaggggn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60

nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnngnttgnt nccttanaag cnaagaagaa 120

ncncanana ancnggcttc taattttttt taaccaccag aaaaatgggn ttttaaantc 180

ggtnggntnc cccggattgg gnccaaggag agagctaaag ttcgttctcg agtctttctg 240

ggatggcaag gaggcagccc gaggatttgc agaaagggtg ctgcttgatc tcaaggatcat 300

ccatctggna agcagatggc tgggtggctgg gatcaaagta ccttccccag caacactttt 360

ctcgttctat gaccaagctt gntcgacgcc accgncaccc tcgggtgccgt cccccccagg 420

tacggntgga ccggcggcga gattggatc gacacctact tctccatggn cagaggtaat 480

gctaccgtgc ctgctatgga gatgaccaat gggtcgacac caa 523

<210> 2688  
 <211> 570  
 <212> nucleic acid  
 <213> Glycine max

<400> 2688

agaggcctgc ttgnctttat canananann ncacngccat gccctccgaa cttccatttc 60

ngcccaggcg tactcccacg cntcctcagt ngccaacga ggctgttcan aangctgctg 120

ctgcattgaa ngnttnggan ggtntcctg caacaaatgt cagggtgccan actggattct 180

caacaaaaga anctcaacct tncaatnctg ncaaccacca ctattggatc cttccctcag 240  
 actgtataac tgaggagggt acgcccctga attnaaagct aacaagatct ccgangaana 300  
 ngtatgttna tgtcanttta aggaggaaat tttcaaantt tgtttgagct tnnataaaag 360  
 cttnatattn atntttctgt catnnagaan nncagatgaa atgntatggt tcnagtncct 420  
 tttngggaca aattttctaa nctttttccn tttacnncnn tatnnggttn gttncaaatc 480  
 ttttgggtcc ccttttctn naaancnncn atcatntttt ngntganttt tatncnccca 540  
 aantnaanna nnccgctttn ttttnttttt 570

<210> 2689  
 <211> 566  
 <212> nucleic acid  
 <213> Glycine max

<400> 2689  
 agaggttnnn ttttttnnat nnaaagtatt tttnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60  
 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnngttagg naaatgacca 120  
 accttagatc tnngaccacc taccngggtg ctttggtat naagggccna antggaggga 180  
 cottgaaaag gctggcatca ctgttatcca aattgatgaa actgctttga gagaaagggt 240  
 ctgccactga gggaaatcaa gaacaaggct cacttacttg gactgggctn gtncatgcct 300  
 ttnagnaatc accaatgttg gnntntccn ngataccact caaaattcac accacatgt 360  
 gctanctcca aacttttaac gacattnttc aatccatnaa ttnacatgg gcccttattg 420  
 ttattcacca tttgaanaac tnttgcttcc gaattaanaa acttcctgt naaagtcntt 480  
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 aaatnccant tccccanaa atnccc 566

<210> 2690  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max

<400> 2690  
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 agcaaggntt tccatctgga aaatttcttt ttgctggtgt tgttgatgga agattnattt 120

gggccaataa ttttgcattct tctctgaaca cccttcaggc acttgagac attgttgaa 180  
atgacaaggt tgtggtttcc acgtcgtgtt ctcttcttca cactgcagtt gatctggtga 240  
atgagaccaa attggaccaa gagattaagt cttggcttgc atttgcagca caaaaagttg 300

<210> 2691  
<211> 239  
<212> nucleic acid  
<213> Glycine max  
<400> 2691

ngtcgcangc acgcgtacgt nagctcgga ttcggctcga gcttgatttn atcaagcaag 60  
gatttccatc tggaaaattt ctntncgctg gtgttggtga tngaaagana tatttggggc 120  
aatagtcttg catcttctct gaacaccctt caggcacttg gggacattgt tgggaatgac 180  
aaggttgtgg ttccacgtc ctgttctctt cttcacactg cagttgatct gtgaatgag 239

<210> 2692  
<211> 303  
<212> nucleic acid  
<213> Glycine max  
<400> 2692

tgcacatcac gcntaagtna gctcggaatt cggctcgagc atttactgcc aatgggtggg 60  
tgcaatcata tggatccgc tgtgtcaaac ctcccatcat ctatgggtgat gtgagccgctc 120  
ccaagcccat gacagttttc tggctcttcaa ctgctcaaag tttgaccaa cgaccaatga 180  
agggaatgct tactggccct gttactattc tgaactggct ctttggttaga gatgaccaac 240  
caagatttga aacatgttac cagattgctt tggctatcaa ggatgagggt gaggatcttg 300  
aga 303

<210> 2693  
<211> 306  
<212> nucleic acid  
<213> Glycine max  
<400> 2693

gtgcacgca cgcgtacgtn agctcggaat tccgctcgag ctgtgtcaag cctcccatca 60  
tctatggtga tgtgagccgt cccaancca tgactgtttn ctggtcttnc aactgctcaa 120

agtttgacca aacgaccaat gaagggaatn cttactggcc ctgttactat tctgaactgg 180  
 tcctttgtta gagatgacca gccaaagattc gaaacatgct accagattgc tttggctatc 240  
 aaggatgagg ttgaggatct tgagaaagca ggtattactg tcatccagat tgatgaagct 300  
 gctctt 306

<210> 2694  
 <211> 459  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2694

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 gaccaagaga ttaagtcttg gcttgcattt gcagcacaaa aagttgttga agtaaatgcc 120  
 ttggccaagg cattgtcttg acagaaggat gaggttttct tttctgctaa tgetgctgcc 180  
 ttggcttcaa ggaagtcctc cccaagggtg ataaatgagg ctgtccaaaa agcogctgct 240  
 gctctgaagg gctctgatca tcggagggcc acaaattgta gtgccagggt ggatgctcaa 300  
 cagaagaaat tgaatctttc tgttcttcca acaactacaa ttggatcttt cctcaaaact 360  
 gccgatctta gaagagttcg tcgtgaattc aaggctaaca agatctccga ggaagattat 420  
 atcccgtttc attaaggagg aaatttacia tgttgtaaa 459

<210> 2695  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2695

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 ctgctgctct gaagggtct gatcatcgga gggccacaaa tggtagtgcc aggttggatg 180  
 ctcaacagaa gaaattgaat ctttctgttc ttccaacaac tacaattggn tctttccctc 240  
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 attata 306

<210> 2696

<211> 285  
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 <213> Glycine max  
  
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 aggaagtcct cccaagggt gataaatgag gctgtccaaa aagccgctgc tgctctgaag 180  
 ggctctgac atcggagggc cacaaatgtt agtgccaggt tggatgctca acagaagaaa 240  
 ttgaatcttt ctgttcttcc aacaactaca attnggtott tccct 285

<210> 2697  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2697  
  
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 cgtgctgct ctgaagggt ctgatcatcg gagggccaca aatgttagtg ccaggctgga 120  
 ttctcaacag aagaaactga atcttctgt tcttccaaca actacaattg ngctctttcc 180  
 ctcaaactgc cgatcttaga agagttcgcc gtgaattcaa ggctaacaag atctccgagg 240  
 aanntatata catttcatta aggagggaaa ttaacaatgt tgtgaagctc caggaagaat 300  
 tga 303

<210> 2698  
 <211> 260  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2698  
  
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 gaccaagaga ttaagtcttg gcttgcatth gcagcacaaa aagttgttga agtaaatgcc 120  
 tggccaaggc attgtctgga agaaggatga ggttttcttt tctgctaattg ctgctgcctt 180  
 ggcttcaagg aagtcctccc caagggtgat aaatgaggct gtccaaaaag ccgctgctgc 240  
 tctgaagggc tctgatcatc 260

<210> 2699  
 <211> 193  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2699  
  
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 ggctgtccaa aaagccgctg ctgctctgaa gggctctgat catcggaggg ccacaaatgt 120  
 tagtgccagg ttggatgctc aacagaagaa attgaatctt tctgttcttc caacaactac 180  
 aattggatct ttc 193

<210> 2700  
 <211> 307  
 <212> nucleic acid  
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 <400> 2700  
  
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 gattgatgag gctgctctaa gagaagggtt accctcgagg aagtctgagg aggttttcta 120  
 tctaaactgg gctgttcact catttaggat taccaactgt ggtgtggagg acactactca 180  
 gattcacact cacatgtgct actccaactt caatgacatc attcactcan tcntagacat 240  
 ggatgctgat gtgatcacca ttgagaactc cagatcagat gagaagttac tttcgggtctt 300  
 ccgtgng 307

<210> 2701  
 <211> 361  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2701  
  
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 ttgctttggg ctatcaagga tgaggttgag gatcttgaga aagcaggtat tactgtcacc 120  
 cagattgatg aagctgctct aagagaaggt ttacctctga ggaagtctga ggaggctttc 180  
 tatctaaact gggctgttca ctcathtaggt attaccaact gtggtgtgga ggacactact 240  
 cagattcaca ctacatgtg ctactccaac ttcaatgaca tcatcactca atcatagaca 300

tggatgctga tgtgatcacc atgagaactc tagatcagac gagaagttac tttcagtctt 360

c 361

<210> 2702

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2702

gtcgcatgca cgcgtacgta agctcggaat tcggctcgag gctgctctaa gagaagggtt 60

acctctgagg aagtctgagg aggcttttcta tctaaactgg gctgttcact catttaggat 120

taccaactgt ggtgtggagg acactactca gattcacact cacatgtgct actccaactt 180

caatgacatc attcactcaa tcatagacat ggatgctgat gtgatcacca ttgagaactc 240

tagatcagac gagaagttac tttcagtctt ccgcgaggga gtgaaatatg gtg 293

<210> 2703

<211> 282

<212> nucleic acid

<213> Glycine max

<400> 2703

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tgaagctgct ctaagagaag gtttacctct gaggaagtct gaggaggctt tctatctaaa 120

ctgggctggt cactcattta ggattaccaa ctgtggtgtg gaggacacta ctcagattca 180

cactcacatg tgctactcca acttcaatga catcattcac tcaatcatag acatggatgc 240

tgatgtgatc accattgaga actctagatc agacgagaag tt 282

<210> 2704

<211> 272

<212> nucleic acid

<213> Glycine max

<400> 2704

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tgaacatgt taccagattg ctttggtat caaggatgag gttgnggatc ttgagaaagc 120

tggtattact gtcattcaga ttgatgaggc tgcnctaaga gaaggtttac ctctgaggaa 180

gtctgaggag gctttctatc taaactgggc tgttactca tttaggatta ccaactgtgg 240  
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<210> 2705  
 <211> 298  
 <212> nucleic acid  
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 cctcccaacc accacaattg gatccttccc tcagactggt gaactgagga ggggtcgctcg 120  
 tgaatacaag gctaacaaga tctcagagga ggagtatggt agttcaatta aagaggaaat 180  
 ccgcaaagtt gttgaactcc aagagaatct tgatatcgat gtcctggtac acggggagcc 240  
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<210> 2706  
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 <212> nucleic acid  
 <213> Glycine max  
 <400> 2706

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 cctcccaacc accacaattg gatccttccc tcagactggt gaactgagga ggggtcgctcg 120  
 tgaatacaag gctaacaaga tctcagagga ggagtatggt agttcaatta aagaggaaat 180  
 ccgcaaagtt gttgaactcc aagagaatct gcatatcgat gtcctggtac acggngagcc 240  
 tgagaggaat gacatggtgg agtactttgg tgagcagttg tcaggctttg cctttaccgt 300  
 taaggct 307

<210> 2707  
 <211> 452  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2707

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 ttctcacttt acgatcaagt actggacaca acagccatgc tcggggcagt tccatctaga 120



tataattgga atggtgggga gattggggtt gatgtttact tctcaatggc aagaggggaat 180  
gcattctgtac cagctatgga aatgaccaag tggtttgaca ccaattacca ttacattggt 240  
cctgaattgg gtctctgatgt taagttctcc tatgcatcac acaaggctgt cgatgaattt 300  
aaagaggcca aagttctggg agttaatact gtacctgtgc ttgtgggacc tgtatcctac 360  
ttgttgctgt caaaaccagc taagggtgtt gaagaagtca tttcccttc tttccctaatt 420  
tgacaagatc cttcctgtct acagggangt tt 452

<210> 2708  
<211> 260  
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<400> 2708

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ttncattac attgttcttg aattgggtcc tgatgttaag ttctcctatg catcacacaa 120  
ggctgtcgat gaatttaaag aggccaaagt tctgggagtt aatactgtac ctgtgcttgt 180  
gggacctgta tctacttgt tgctgtcaaa accagctaag ggtgttgaga agtcattttc 240  
ccttctttcc ctaattgaca 260

<210> 2709  
<211> 275  
<212> nucleic acid  
<213> Glycine max

<400> 2709

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gttccatcta gatataattg gaatgggtgg gagattgggt ttgatgttta cttctcgatg 120  
gcaagaggga attcatctgt accagctatg gaaatgacca agtggtttga cacaaactat 180  
cattacattg ttctgaatt ggggccagat gttaagttct cctatnccat cacacaagggt 240  
tgtggatgaa tacntngagg ctaaagttct gggaa 275

<210> 2710  
<211> 304  
<212> nucleic acid  
<213> Glycine max

<400> 2710

ntnacgcnta cgtaagctcg gaattcggct cgagctagtc tcctactctc acccgcaaga 60  
aaaaaaatgg catctcacat cgttggatac ccccgatgg gtcccaagan agagctcaag 120  
ttcgctctcg agtctttctg ggatggcaag agcagcgccg aggatttgca gaaggtgtct 180  
tctgatctca gggcatccat ctggaagcag atggctgatg ctgggatcaa gtacatcccc 240  
agcaacactt tctctcacta tgaccagggt ctcgacgcca ccgccaccct cggtgccggt 300  
ccac 304

<210> 2711

<211> 341

<212> nucleic acid

<213> Glycine max

<400> 2711

tcgcangcac gcgtacgtaa gctcggaatt cggtcgcagc tccactcaga agcgaagaan 60  
aagccacaga gaactagtct cctacttctc acccgcaaga naaaaaatggc atctcacatc 120  
gttggatacc cccgatggg tcccaagaga gagtcaagt tcgctctcga gtctttctgg 180  
gatggcaaga ncagcgccga ggatttgagc aaggtgtctt ctgatctcag ggcatccatc 240  
tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaacacttt ctctcactat 300  
gaccagggtt cgcgacccac cgccaccctc ggtgccgttc c 341

<210> 2712

<211> 322

<212> nucleic acid

<213> Glycine max

<400> 2712

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gaaccagtct cctactctct ctcaccacca agaaaaatgg catctcacat cgttggatac 120  
cccgcatgg gtcccaagag agagctcaag ttcgctctcg agtctttctg ggatggcaag 180  
agcagcgccg aggatttgca gaaggtggct gctgatctca ggcatccat ctggaagcag 240  
atggctggtg ctgggatcaa gtacatcccc agcaacactt tctcgttcta tgaccagctg 300  
ctcgacgcca ccgccaccct cg 322

<210> 2713  
 <211> 328  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2713  
  
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 agttcgctct cgagtctttc tgggatggca agagcagcgc cgaggatttc agaaggtgtc 180  
 ttctgatctc agggcatcca tctggaagca gatggctgat gctgggatca agtacatccc 240  
 cagcaacaact ttctctcact atgacanggt tctcgacgcc accggcacc cgggtgccgt 300  
 tncaccaagt aggtggancc gcggcgag 328

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 <213> Glycine max  
  
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 catctcacat cgttggatac ccccgcatgg gtcccaagag agagctcaag ttcgctctcg 180  
 agtctttctg ggatggcaag agcagcgccg aggatttgca gaagggtggct gctgatctca 240  
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 tctcgttcta tgaccagctg ctcgacgcca ccgccaccc 339

<210> 2715  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2715  
  
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 tctctcacc acaagaaaaa tggcatctca catcggttga taccctcgca tgggtcccaa 120  
 gagagagctc aagttcgctc tcgagctctt ctgggatggc aagagcagcg ccgaggattt 180

gcagaaggtg gctgctgac tcaggtcatc catctggaag cagatggctg gtgctgggat 240  
 caagtacatc cccagcaaca ctttctcggt ctatgaccag ctgctcgacg ccaccg 296

<210> 2716  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2716

ngtngcangc acgcgtacgt aagctcgga ttcggctcga gtttgagaa ggtgtcttct 60  
 gatctcaggg catccatctg gaagcagatg gctgatgctg ggatcaagta catccccagc 120  
 aacactttct ctactatga ccaggttctc gagccaccg ccaccctcgg tgcggtccac 180  
 caaggtaagg ctggcaccgg cggcgagatt gggtttgata cctacttcnc catggccaga 240  
 ggtaatgcta ccgtgccagc tatggagatg accaagtggg tcgacaccaa ctaccacnct 300  
 nttgtccc 308

<210> 2717  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2717

anntnnnctg acgtaanctc ggaattcggc tcgagcttgc tccctcanan ncntgaanaa 60  
 gccacagaga actagtctcc taactctcac ccgcaagacn aaaaatggca tctcacatcg 120  
 ttggnatccc ccgtatnggt cccaagagag agcncaagtt cgctctcgag tctttctggg 180  
 atggcaagag cagcgccgag gatttgaga aggtgtcttc tgatctcagg gcatccatct 240  
 ggaagcagat ggctgatgct gggatcaagt acatccccag caacactttc tctcactatg 300  
 accaggttct cgac 314

<210> 2718  
 <211> 307  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2718

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctccctcaga agcgaagaag 60

aagccacaga gaaccagtct cctactctct ctcacccaca agaaaaatgg catctcacat 120  
 cgttggatac ccccgcatgg gtcccaagan agagctcaag ttcgctctcg agtctttctg 180  
 ggatggcaag agcagcgccg aggatttgca gaaggtggct gctgatctca ggcatccat 240  
 ctggaagcag atggctgggtg ctgggatcaa gtacatcccc agcaacactt tctcgttcta 300  
 tgaccag 307

<210> 2719  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2719

acgcgtacgt aagctcggaa ttcggctcga ggcttcaact tgctccctca gaagcgaaga 60  
 agaagccaca gagaaccagt ctctactctt ctctcaccca caagaaaaat ggcatctcac 120  
 atcgttggat acccccgcat ggggtcccaag agagagctca agttcgctct cgagtctttc 180  
 tgggatggca agagcagcgc cgaggatttg cagaaggtgg ctgctgatct caggatccatcc 240  
 atctggaagc agatggctgg tgctgggatc aagtacatcc ccagcaaacac tttctcgttc 300  
 tatgaccact gctcgacg 318

<210> 2720  
 <211> 324  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2720

gcacgcgtac gtaagctcgg aattcggtc gagctcagaa nccaanaaga agccacncna 60  
 gaactagtct cctactctca cccgcaanaa aaaaatggca tctcacatcg ttngatacc 120  
 cccgtatggg tccaagaga gagctcaagt tcgctctcga gtctttcttg gatggcaaga 180  
 gcagcgccga ggatttgca aaggtgtctt ctgatctcag ggcatccatc tggaagcaga 240  
 tggctgatgc tgggatcaag tacatcccca gcaaacactt ctctcactat gaccagggtc 300  
 tcgacgccac cgcaccctc ggtn 324

<210> 2721  
 <211> 331  
 <212> nucleic acid

<213> Glycine max

<400> 2721

nnncnngca tncgtacgta agctcggaat tcggctcgag cggctcgagt tcaacttget 60  
ccctcagaag cgaagaagaa gccacagaga actagtctcc tactctcacc cgcaagaaaa 120  
aaatggcatc tcacatcggtt ggataccccc gtatgggtcc caagagagag ctcaagtctg 180  
ctctcgagtc tttctgggat ggcaagagca gcgcgagga tttgcagnag gtgtcttctg 240  
atctcagggc atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 300  
acactttctc tcactatgac caggttctcg n 331

<210> 2722

<211> 327

<212> nucleic acid

<213> Glycine max

<400> 2722

gttgcangca cgcgtacgta agctcngaatt tcggctcgag gccacagaga ctccagtctc 60  
ctactctctc tcaccanaa gananntggn atntcacntc gttggatacc cccgcatggg 120  
tcccaagana gagctcnagt tcgctctcga gtctttctgg gatggcnagn nongcgccga 180  
ggatttggca naaggtggct gctgatctca ggtcatccat ctogaagcag atggctggtg 240  
ctgggatcaa gtacntcccc agcaacactt tctcgttcta tgaccagctg ctgacgccca 300  
ccgccaccct cggtgccgtc cccccag 327

<210> 2723

<211> 316

<212> nucleic acid

<213> Glycine max

<400> 2723

gtcgcattgca cgcgtacgta agctcggaat tcggctcgag ggctgatgct gggatcaagt 60  
acatccccag caacactttc tctcactang accnggtttc tcgacgccac cgcnacccctc 120  
ggtgccgttc caccaaggna cggctggacc ggcggcgaga ttgggtttga tacctacttc 180  
tccatggcca nangtaatgc taccgtgcca gctatggaga tgaccaagtg gttcganacc 240  
aactancact ttattgtccc tgaatgggcn ctgatgtgaa cttcacctat gcttctcaca 300

aggcngctga tgaana

316

<210> 2724  
<211> 320  
<212> nucleic acid  
<213> Glycine max

<400> 2724

cgntnnnaag cgtacgttag ctcggaattc ggctcgagct tgctccctca gaagcgaaga 60

agaagccaca gagaaccagt ctctactct ctctaccca caagaaaaat ggcatctcac 120

atcggttgat acccccgcat ggggcccaag agagagctca agttcgctct cgagtcttct 180

gggatggcaa gaggcgcgc gaggatttgc agaaggtggc tgctgatctc aggtcatcca 240

tctggaagca gatggctggg gctgggatca agtacatccc cagcaacact ttctcgttct 300

atgaccagct gctcgacgcc 320

<210> 2725  
<211> 301  
<212> nucleic acid  
<213> Glycine max

<400> 2725

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctgagaagcg aagaagaagc 60

cacagagaac tagtctcta ctctacccg caagaaaaaa atggcatctc acatcgttgg 120

atacccccgt atgggtccca agagagagct caagttcgct ctcgagtctt tctgggatgg 180

caagagcagc gccgaggatt tgcagaaggt gtcttctgat ctgagggcat ccatctggaa 240

gcagatggct gatgctggga tcaagtacat cccagcaac actttctctc actatgacca 300

g 301

<210> 2726  
<211> 312  
<212> nucleic acid  
<213> Glycine max

<400> 2726

gtngcatgcn cgcgtacgta agctcggnat tcggctcgan ctgagccga atcggctcga 60

gagaaaaatg gcattctaca tcgttagata ccccgcatg ggncccaaga gagagctcaa 120

gttcgctctc gagtctttct gggatggcaa gaggcagcc gaggtttgc agaaggtggc 180  
 tgotgatctc aggtcatcca tctggaagca gatggctggt gctgggatca agtacatccc 240  
 cagcaacact ttctcgttct atgaccagct gctnccgacgc caccgccacc ctgggtgccc 300  
 tccccccan gt 312

<210> 2727  
 <211> 301  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2727

nnogtctcan gcacgcgtac gtnagctcnn naattcggct cgaggaacta gtctcctact 60  
 ctcanccnca ananaataat ggcacatctgca catgcgntgg atacccccgt atgggtccca 120  
 agagnnanct caagttcgct ctcgagtctt tctgggatgg caagagcacc gccgaggatt 180  
 tgcagaaggt gtcttctgat ctccagggcat ccacatctggaa gcagatngct gatgctggga 240  
 tcaagtacat cccanacaac actttctctc actatgacca ggtncctgac gccnccgcca 300  
 t 301

<210> 2728  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2728

angtcgcang cagcgcgtacg taagctcgga attcggctcg agtgcttcaa ctgctccct 60  
 cagaagcgaa gaagaagcca cagagaacca gtctcctact ctctctcacc cacaagaaaa 120  
 atggcatctc acatcggttg atacccccgc atgggtccca agagagagct caagttcgct 180  
 ctcgagtctt tctgggatgg caagagcagc gccgaggatt tgcagaaggt ggctgctgat 240  
 ctccaggtcat ccacatctggaa gcagatggct ggtgctggga tcaagtacat cccagcaac 300  
 actttctcgt tctatg 316

<210> 2729  
 <211> 330  
 <212> nucleic acid  
 <213> Glycine max



<400> 2729

natctcangc aagcgtacgt attctcgnna ttnggcnacg aggcttcaac ttgctccctc 60

agaancgaag aagaagnnac agananctan tctcctactc tcacccgcaa gaanaaaatn 120

gcattctcaca tcgttgata ccccgatatg ggtcccaaga gagagctcaa gttcgctctc 180

gagtctttct gggatggcaa gagcagcgn gaggatttgc agaaggtgtc ttctgatctc 240

agggcattcca tctggaagca gatggctgat gctgggatca agtacatccc cagcaaacact 300

ttctctcata tgcaccaggt tctcgaagcc 330

<210> 2730

<211> 271

<212> nucleic acid

<213> Glycine max

<400> 2730

cccgatatggg tccaagaga gagctcaagt tcgctctcga gtctttcttg gatggcaaga 60

gcagcgccga ggatttgcag aaggtgtctt ctgatctcag ggcattccatc tggaagcaga 120

ggctgatgct gggatcaagt acatccccag caacactttc tctcactatg accaggttct 180

ngacgcnacc gncannctcg gtgntgtnc c atnaaggnan gnnatatann gttgtnatnn 240

nggntnnann ntnttttcca ngttntgcag g 271

<210> 2731

<211> 318

<212> nucleic acid

<213> Glycine max

<400> 2731

tcccgtagct nagctcgga ttcggctcgn ggcttcanct tgctccctca gaagcgaaga 60

agaagccaca gagaactagt ctctactcn canccncnag anaaaaatgg catctcacat 120

cgttgatgata ccccgatatg gtcccaagan agagctcaag ttcgctctcg agtctttctg 180

ggatgcaaga gcagcgccga ggatttgcag aaggtgtctt ctgatctcag ggcattccatc 240

tggaagcaga tggctgatgc tgggatcaag tacatcccca gcaaacacttt ctctcactat 300

gaccaggttc tcgacgca 318

<210> 2732

<211> 307  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2732  
  
 acangcatgc acnggtacgt aagctcggaa ttcggtctga gctctgcttc aacttgetcc 60  
 ctcagaagcg aagaagaagc cacagagaac tagtctccta cttcttcacc cgcaaganaa 120  
 aaatggcatc tcacatcggt ggataacccc gtatgggtcc caaganagag ctcaacttcg 180  
 ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagaag gtgtcttctg 240  
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 acacttt 307

<210> 2733  
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 <212> nucleic acid  
 <213> Glycine max  
  
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 gaagcgaaga agaagccaca gagaaccagt ctctactac tactctcacc cacaagaaaa 120  
 atggcatctc acatcggttg atacccccgc atgggtccca agagagagct caagtctgct 180  
 ctcgagtctt tctgggatgg caagagcagc gccgaggatt tgcagaaggt ggctgctgat 240  
 ctcaggtcat ccatctggaa gcagatggct ggtgctggga tcaagtacat ccccagcaac 300  
 actt 304

<210> 2734  
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 acccacaaga naaatgncat ctcacatgcn tggatanccc cgcattgggtc ccnaganaga 180  
 gctcaagttc gcnctcgagt ctttctggga tggcaagagc agcgccgagg atttgcagaa 240

ggtaggtgct gatctcaggt catccatctg gaagcagatg gctggtagct ggatcaagta 300  
catccccagc aacactttct cgttctatga cca 333

<210> 2735  
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<212> nucleic acid  
<213> Glycine max  
  
<400> 2735

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agaagaagcc acagagaact agtctcctac tctcaccgcg aagaaaaaaaa tggcatctca 120  
catcgttgga tcccccgta tgggtcccaa gagagagctc aagttcgctc tcgagctctt 180  
ctgggatggc aagagcagcg ccgaggattt gcagaagggt tcttctgata tcagggcatt 240  
catctggaag cagatggctg atgctgggat caagtacatc ccagcaaca cttctctc 299

<210> 2736  
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<212> nucleic acid  
<213> Glycine max  
  
<400> 2736

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gccacagaga actagtctgc ctactctcac ccgcaagana aaaatggcat ctacatgcg 120  
ttggataccc ccgtatgggt cccaagagag agctcaagtt cgctctcgag tctttctggg 180  
atggcaagag cagcgccgag gatttgaga aggtgtcttc tgatctcagg gcatccatct 240  
ggaagcagat ggctgatgct gggatcaagt acatccccag caacactttc tctactatg 300  
accaggttct tcgacgcnac gccacctcgg tgc 333

<210> 2737  
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<212> nucleic acid  
<213> Glycine max  
  
<400> 2737

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agaagaagcc acagagaact agtactcnct antctcaccg gcaagannna aatggcatct 120

cacatcgntg gatacccccg tatgggtccc aagagagagc tcnagttcgt nctcgagtct 180  
 ttctgggatg gcaagagcag cgccgaggat ttgcagaagg tgtcttctga tctcagggca 240  
 tccatctgga agcagatggc tgntgctggg atcaagtaca tccccagcaa cactttctct 300  
 nactatggcc agttctcnac 320

<210> 2738  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max  
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 gaagccacag agaaccagtc tcctacttct ctctcaccca caagaaaaat ggcatctcac 120  
 atcgttggat acccccgcat ggggtccaag agagagctca agttcgtctt cgagtctttc 180  
 tgggatggca agagcagcgc cgaggatttg cagaaggtgg ctgctgatct caggtcatcc 240  
 atctggaagc agatggctgg tgctgggatc aagtacatcc ccagcaa 287

<210> 2739  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
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gttgcattgn agcgtacgta agcttcggaa ttctggtnnn ggatttggat cgagcttntt 60  
 ccctcagnag cgnagaagaa gccncagaga actagtctcn tactctcacc cgcaagaaaa 120  
 aaatggcatc tcacatcgtt ggatacccc gtatgggtcc caagagagag ctcaagttcg 180  
 ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagaag gtgtcttctg 240  
 atctcagggc atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca 300  
 acactt 306

<210> 2740  
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 <212> nucleic acid  
 <213> Glycine max  
 <400> 2740

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tcacatcggt ggataccccc gcatgggtcc caagagagag ctcaagttcg ctctcgagtc 180  
tttctgggat ggcaagagca gcgccgagga tttgcagaag gtggctgctg atctcaggtc 240  
atccatctgg aagcagatgg ctggtgctgg gatcaagtac atccccagca a 291

<210> 2741  
<211> 322  
<212> nucleic acid  
<213> Glycine max  
<400> 2741

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agccacagag aactagtctc ctactctcac ccgcaagaaa aaaatggcat actgcacatt 120  
cgttggatac ccccgatagg gtcccaagan agagctcaag ttcgnnctcg agtctttctg 180  
ggatggcaag cgcagcgccg aggatttgca gaaggtgtct tctgatctca gggcatccat 240  
ctggaagcag atggctgatg ctgggatcaa gtacatcccc agcaacactt tctctcacta 300  
tgaccagttc tcgacgccac gn 322

<210> 2742  
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<212> nucleic acid  
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<400> 2742

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agccacagag aactagtctc ctactctaca cccgcaagaa aaaaatggca tctacacatc 120  
gttggatacc cccgataggg tcccaagaga gagctcaagt tcgctctcga gtctttctgg 180  
gatggcaaga gcagcgccga ggatttgcag aaggtgtctt ctgatctcag ggcattccatc 240  
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accaggttct 310

<210> 2743  
<211> 304  
<212> nucleic acid

<213> Glycine max

<400> 2743

agttgcangc acgcgtacnt aagctcggaa ttcggctcga gtcagaagcg aagaagaagc 60  
cacagagaac tagtcnnnt actctcacc cgaagaaana aatngccatc tcanatgcgt 120  
tggatncccc cgtatgggtc ccaagagaga gctcaagttc gctctcgagt ctttctggga 180  
tggcaangnc ancgccgagg atttgcagaa ggtgttttct gatctcaggg catccatctg 240  
gaagcagatg gctgatgctg ggatcaagta catccccagc aacactttct ntcactatga 300  
ccag 304

<210> 2744

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2744

tcgcangnac nntcggaaat tcggctcgag cttgctccct cagaagcgaa gaagaagcca 60  
cagagaacca gtctcctact ctctctcacc cacaagaaaa atggcatctc acatcggttg 120  
atacccccgc atgggtccca agagagagct caagttcgct ctcgagtctt tctgggatgg 180  
caagagcagc gccgaggatt tgcagaaggt ggctgctgat ctcaggatcat ccactctggaa 240  
gcagatggct ggtgctggga tcaagtacat cccagc 277

<210> 2745

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2745

gtngcangca cgcgtacgta agctcggaaat tcggctcgag ctccctcaga agcgaagaag 60  
aagccacaga gaactagtct cctactctca cccgcaagan aaaaatggca tctcacatcg 120  
ttggataccc ccgtatgggt cccaagagng agctcaagtt cgctctcgag tctttctggg 180  
atggcaagag cagcgccgag gatttgcaga aggtgtcttc tgatctcagg gcatccatct 240  
ggaagcagat ggctgatgct gggatcaagt acatccccag caacactt 288

<210> 2746

<211> 318  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2746  
  
 gttgcangca cgcgtagcta agctcgggaat tcggctcgag cagaagcgaa gaagaagcca 60  
 nagagaacta gtctcctaen nttcaccgcg aagaaaaaat ggnatctcac atcgttggat 120  
 acncccgat gggtnnccaa gagagngnna agttcgntct cgagtccttc tgggatggca 180  
 agagcagcgc cgaggatttg cagaagggtg cttctgatct cagggcatcc atctggacgc 240  
 agatggctga tgctgggacg aagtacatnn ncagcaanac tttctctcan tatgaccagg 300  
 ttctcgacgc naccgcca 318

<210> 2747  
 <211> 331  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2747  
  
 gcaaantcgc tgnangenta cgtnagctcg ganttcggat ngagctcgag ccgctgcttc 60  
 aacttgctac ctcagaagcg aagaagaagc cacagagaac cagtctccta ctctctctca 120  
 cccacaagaa naatgggacg tcacatcggt ggataccccc gcatgggtcc caagagagag 180  
 ntcnagttcg ctctcgagtc tttctgggat ggcaagngcn gcgccgagga tttgcagaag 240  
 gtggctgctg atctcaggtc atccatctgg aagccagatg gctgggtgctg ggatcaagta 300  
 catccccagc aacactttct cgtttatgnc c 331

<210> 2748  
 <211> 307  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2748  
  
 aatcgcangc aagcgtacgt aagctcgga ttcggctoga gctctgcttc aacttgctcc 60  
 ctcagaagcg aagaagaagc cacagagaac cagtctccta ctctctctca cccacaagaa 120  
 aaatggcatc tcacatcggt ggataccccc gcatgggtcc caagagaagc tcaagttcgc 180  
 tctcgagtct ttctgggatg gcaagagcag cgccgaggat ttgcagaagg tggctgctga 240

tctcagggtca tccatctgga agcagatggc tgggtgctggg atcaagtaca tccccagcaa 300  
cattttct 307

<210> 2749  
<211> 302  
<212> nucleic acid  
<213> Glycine max  
<400> 2749

gtcgcangna cgcgtacgta agctcggaat tcggctcgag cttgntccct cagaagcgaa 60  
gaanaancca cagagaanta gtctnctact ctgcacccgc aanaaaaaaa tggcntctca 120  
catgcgttgg atacccccgt atgggtccca aganagagct caagttcgct ctcgagtctt 180  
tctgggatgg caagagcagc gcgaggatt tgcagaaggt gtcttctgat ctcagggcat 240  
ccatctggaa gcagatggct gatgctggga tcaagtacat cccagcaac actttctctc 300  
an 302

<210> 2750  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
<400> 2750

cangcacgcg tacgtnagct cggaattcgg ctcgagnctt gctccctcag aagcgaagaa 60  
gaagccacag agaaccagtc tcctactctc tctcaccac aaganaaatg gcattctcaca 120  
tcgttgata ccccgcatg ggtcccaaga nagagctcaa gtctgctctc gagtncttct 180  
gggatggcaa gacgagcgcc gaggatttgc agaaggtggc tntgatctc aggtcatcca 240  
tctggnagca gatggctggg gctgggatca agtacatccc cagcaac 287

<210> 2751  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2751

agtncangc acgcgtacgt aagctcgga ttcggctcga gctggcttca anttgctccc 60  
tcagaagcga agaagaagcc acagaganct ngctctctac tctcaccgc aaganaaaaa 120



tggcatctca catcgttggga tcccccgta tgggtcccaa ganagagctc aagttcgctc 180  
 tcgagtcttt ctgggatggc aagagcagcg ccgaggnttt gcagaagggtg tcttctgntc 240  
 tcagggcata catctggaag cagatggctg atgctgggat caagtacatc cccagcaaca 300

<210> 2752  
 <211> 285  
 <212> nucleic acid  
 <213> Glycine max

<400> 2752

ncttnacgc acgcgtacgt nagctcggaa ttcggctcgg gctccctcag aagcgaagaa 60  
 gaagccacag agaaccagtc tcctantcnc tctcaccac aagaaaaatg gcattctaca 120  
 tcgttggata cccccgcatg ggtcccaaga nagagctcaa gttcgctctc gagtctttct 180  
 gggatggcaa gagnacgcc gaggatttgc agaagggtggc tgctgatctc aggtcatcca 240  
 tctggaagca gatggctggg gctgggatca agtacatcnc cagcn 285

<210> 2753  
 <211> 326  
 <212> nucleic acid  
 <213> Glycine max

<400> 2753

tgctgtaana nanaangtng cgtgcacgcn tacgtaaagc tcgggaattc ggctcgagct 60  
 ggnttcaant tggntcctca naagcgaaga agaagccaca gagaaccagt ctctantnt 120  
 ctntcaccca caagaaaaat ggcntctcac atcgttggat acccccgcat ggggtcccaag 180  
 anagagtca agttcgntnt cgagtctttc tgggatggca agancagcgc cgaggatttg 240  
 cagaagggtg ctgntgatnt caggtcatcc atntggaagc agatggctgg tgntgggatc 300  
 aagtacatcc ccagcaanac tttntc 326

<210> 2754  
 <211> 281  
 <212> nucleic acid  
 <213> Glycine max

<400> 2754

ncgtcgcang cacgcgtacg taagctcgga attcggctcg aggtccctc agaagcgaag 60

aagaagccac agagaaccag tctctactc tctctcacc acaagaaaaa tggcatctca 120  
catcggttga taccocccgca tgggtcccaa gagagagctc aagttcgctc tcgagtcctt 180  
ctgggatggc aagagcagcg ccgaggattt gcagaaggtg gctgctgac tcaggtcatc 240  
catctggaag cagatggctg gtgctgggat caagtacatc c 281

<210> 2755  
<211> 303  
<212> nucleic acid  
<213> Glycine max  
<400> 2755

gtcgcangca cgcgtncgac gantaogtna nctcggnntt nggntccncg ctnacatggg 60  
catngnaaac tgtttacgag tgagtntcct angencancc gcaagaaaaa aatggcatct 120  
cacatcggtg gatacccccg natgggttcc caagagagag ctcaagttcg ctctcgagtc 180  
tttctgggat ggcaagagca gcgccgagga tttgcagaag gtgtcttctg atctcagggc 240  
atccatctgg aagcagatgg ctgatgctgg gatcaagtac atccccagca acatttctct 300  
cat 303

<210> 2756  
<211> 300  
<212> nucleic acid  
<213> Glycine max  
<400> 2756

cgtcgcatgc acgcgtacgt nagctcggnn ttcggctcgn gcggctcng gcttcaactt 60  
gctccctcag aagcgaagaa gaagccacag agaactagtc tctaacctc acccgcaaga 120  
naaaaatggc atctcacatc gttggatacc cccgtatggg tcccaagaga gagctcaagt 180  
tcgtctcga gtctttcttg gatggcaaga gcagcgccga ggatttgcag aaggtgtctt 240  
ctgatctcag ggcattccatc tggaagcaga tggctgatgc tgggatcaag tacatcccca 300

<210> 2757  
<211> 299  
<212> nucleic acid  
<213> Glycine max  
<400> 2757

tngcangcac gcgtacgtaa gctcgggaatt cggctcgagc tgcttcaact tgctccctca 60  
gaagcgaaga agaagcnaca gaggnccagt ctctactct ctctcaccca caagaaaaat 120  
ggcatctcac atcgttggat acccccgcat gggteccaag agagagctca agttcgctct 180  
cgagtctttc tgggatggcn ngagcagcgc cgaggatntg cagaaggtgg ctntgatct 240  
caggncatcc atctggaagc aganggctgg tgcngggatc aagtacatcc ncagcnacn 299

<210> 2758  
<211> 310  
<212> nucleic acid  
<213> Glycine max  
<400> 2758

tnncatgcac gcgtacgtaa gctcgggaatt cggctcgagg ctccctcnga agcgaagaag 60  
aagccacaga gaactagtcn nctantotca cccgcaagaa naaaatgngc atcttcacat 120  
ncgttggata ccccgctatg ggtcccaaga gaganctcaa gttecgctctc gagtctttct 180  
gggatggcaa gaggcagcnc gaggatttgc agaaggtgtc ttctgatctc agggcatcca 240  
tctggaagca gatggctgat gctgggatca agtacatccc cagcaacact ttctnccan 300  
tntgaccagg 310

<210> 2759  
<211> 318  
<212> nucleic acid  
<213> Glycine max  
<400> 2759

acgcatgnag cgtacgnag ctccgnattc ggctcgagct gcttcaactt nctccctcag 60  
aagcgaagaa gaagccacag agaactantc tctactctc acccgacta aaanaatggc 120  
atctcacatc gttggatacc cccgtatngg ncccaagana gagctcaagt tcgctctcga 180  
gtctttcttg gatngcnnga ncagngccgn ggatttgcac caggtgtctt ctgatctcan 240  
ggcatccatc tggnnacaga tggctgntgc tgggatcaag tncatccnca ncaacacttt 300  
ctctcactag nncaggtt 318

<210> 2760  
<211> 293  
<212> nucleic acid

<213> Glycine max

<400> 2760

nncgtcgcan gcacgcgtac gtaagctcgg taattcggct cgagnaactt gctccctcag 60  
aagcgaagna gaagccacag agnactagtc tcctacttct caccgcgnag anaaanntgg 120  
catctcacat cgttggatac ccccgatatg gtcccaagag agagctcaag tncnctctcn 180  
agtctttctg ggatggcnag agcagcgccg aggatttgca gaaggtgtct nctnatctca 240  
gggcatccat ctggaagcng atggctgatg ctgggatcaa gtacatcccc agc 293

<210> 2761

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 2761

ngntcangca cncgtacgtn agctcggaat tcnctcgan gctccctcag aagcgaagaa 60  
nnagccacag agaactngnc tcctactctc acccgcaaga aaaaaatggc acctcanntc 120  
gttggatacc cccgtatgng tcccaagagg gagctcaagt tcgctctcga gtcttctggg 180  
atggcaagag cagcgccgag gatttgca gaaggtgtctc tgatctcagg ggcattccatc 240  
tggaagcaga tggctgatgc tgggatcaag tacatcccc gcancacttt ctctcactat 300  
gn 302

<210> 2762

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2762

gatgctngca cgcgtacgtn agctcggaat tnggctcgag cagagaacta gtctcctact 60  
ctcaccgcga agaaaaaat ngnatctcac atcgttggat acccccgat ggggtcccaag 120  
agagagcnca agttcgctct cgagttcttt ctnggatggc aagngcagct ccgaggattt 180  
gcagaaggtg tcnnntgatc tcagggcctc catctggaag cagatggctg atgctgggat 240  
caagtacatc cccngcaaca ntttctcnna ctctgacaa ggttctcgac genaccgcga 300  
accctcgggtg n 311

<210> 2763  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2763  
  
 cncgtacgta agctcggaat tcggctcgag cnaagaanaa gccacagcag anccagtctc 60  
 ctantctctc tcanccacaa nanaaatngc atencacatc gtnggatacc cccgcatggn 120  
 taccceanag agagnncaag ttacgctctc gagtctttct gggatggcaa gagcagcgcc 180  
 gaggatttgc agaaggtggc tgctgntctc aggtcatcctn tctggaagca gatggctggt 240  
 gctgggatca agtacatccc cancaacact ttctcgttct atgaccagct gctcgacg 298

<210> 2764  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2764  
  
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 gaagcgaaga agaagccaca gagaaccagt ctctactctc ctctaccca caagaaaaat 120  
 ggnatctcac atcgttggat acccccgcac gggncccaag agagagctcn agttcgntct 180  
 cgagtctttc tgggatggna anagcancgn cgangntttg canaangngg ctgctggtct 240  
 cangncatcc atctggaanc ngatggctgg tgctgggatc nagtacatcc ccagcaacac 300

<210> 2765  
 <211> 330  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2765  
  
 gtgcgatgca cgcgtacgta agctcggaat tcggctcgag gcttcaactt gctccctcag 60  
 aagcgaagaa gaagccacag agaactagtc tccttacttc tcacccgcaa ganaaaaaatg 120  
 ggcatctcac atcgttggat acccccgtat gggcccaag anagagctca agttcgctct 180  
 cgagtctttc tgggatggca agagcagcgc cgaggatttg cagaagggtg cttctgatct 240  
 caggcatcc atctggaagc agatggctga tgctgggatc aatacatccc cagcaacatt 300

tctctcatat gaccagttct cggacgccac

330

<210> 2766

<211> 308

<212> nucleic acid

<213> Glycine max

<400> 2766

ttgangcacg cgtacgtnag ctcggaattc ggctcgagct tcaacttgct ccctcagang 60

cgaagaagan gccacagaga acnagtctcc tactctcacc cgcaaganaa aaatggcatc 120

tccacatcgt tggatacccc cgtatgggtn cccaaganag agctcaagtt cgctctcgag 180

tctttctggg atggcaagag cagcgccgag gatttgcaga aggtgtcttc tgatctcagg 240

gcatccatct ggaagcagat ggctgatgct gngantcaag tacatcccca gcaacacttt 300

ctctcact 308

<210> 2767

<211> 309

<212> nucleic acid

<213> Glycine max

<400> 2767

gttnaatgca cgcgtacgta agctcggaat tcggctcgag gctccctcag aagcgaagaa 60

gaagccacag agaaccagtc tctactctc tctcaccac aanaaaaatg gcatctcaca 120

tcgttggata cccccgatg ggtcccaaga gagagctcaa gtctgctctc gactctttct 180

gggatggcaa gagcagcgc gaggatttgc agaagggtggc tgctgatctc aggttcatcc 240

atctggaagc agatggctgg tgctggggat caagtacatc cccagcaaca cttctcgttc 300

tatgaccag 309

<210> 2768

<211> 247

<212> nucleic acid

<213> Glycine max

<400> 2768

ctanaactnn acnccctcag aagcgaagaa gaagccacag agaactagtc tctactctc 60

acccgcaaga naaaaatggc atctcacatc gttggatacc cccgtatggg tccaagaga 120

gagctcaagt tcgtctcgag tcttctggg atggcaagag cagcgccgag gatttgcaga 180  
 aggtgtcttc tgatctcagg gcatccatct ggaagcagat ggctgangct gggntcaagt 240  
 acatccc 247

<210> 2769  
 <211> 248  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2769

gcacgcgtac gtaagctcgg aattcggctc gagctccctc agaagcgaag aanaagccac 60  
 agagaactag tctnctactc tcannccgca agaaaaaaat ggcattctcac atcgttggat 120  
 acccccgtat ggggtcccaag agagagctca agtctcgtct cgagtctttc tgggatggca 180  
 agagcagcgc cgaggatttg cagaaggtgt cttctgatct cagggcatcc atctggaagc 240  
 agatggct 248

<210> 2770  
 <211> 284  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2770

ncgttgcang cagcgcgtacg tnagctcgga attnngctcg agtctgcttc aacttgctcc 60  
 ctcagaancg aagaagaagc cacagaganc tagtctccta ctctcaccgc caagaaaaaa 120  
 atggcatctc acatcgttgg ntacccccgt atgggtccca agaganagct caagtctgct 180  
 ctcgagtctt tctgggatgg cnagngcagc gccgaggatt tgcagaaggt gtcttctgat 240  
 ctcagggcat ccattctggaa gcagatggct gangctggga tcan 284

<210> 2771  
 <211> 329  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2771

anaancaatg caccgtacnt aantcggntc natgncgagc ngaattcggc tcgagctctg 60  
 cttcaacttg ctcccatnga agcgaanaag aagccacaga nnactantct cctantctca 120

cncgcaagan aanaatggca nctcacatgc gttggatacc ccntatggg tcccaagaga 180  
gagctcaagt tcgctctcga gtctttctgg gatngcaaga gnagcgccga ggatttgcn 240  
aaggngtctt ctgatctcag ggcattccatc tggnagcana tggctgatgc tnggancaag 300  
tacatcccca gcaacatttc tctcatagn 329

<210> 2772  
<211> 302  
<212> nucleic acid  
<213> Glycine max  
<400> 2772

tgncgcntac ntaagnncgg aattcggctc gnggcgaaga agaagccaca gagaaccagt 60  
ctcctactct ctctaccca caagaaaaat ggcattctac atgcgttgga tcccccgca 120  
tggttaccca agagagagct caagtctcgt ctgagctctt tctgggatgg caagagcagc 180  
gccgaggatt tgcagaaggt ggctgctgat ctgaggtcat ccatnctgga agcagatggc 240  
tggtgctggg atcaagtaca tcccnagcaa cacttctcgn tctatgacca gctgcnnacg 300  
cc 302

<210> 2773  
<211> 281  
<212> nucleic acid  
<213> Glycine max  
<400> 2773

gcacgcgtac gtaagctcgg aattcggctc ganctcgagc cgattcggct cgagttgctc 60  
cctcagaagc gaagaaacng ccacagagaa ccagtctcct actctctctc aaccacaaga 120  
aaaatggcat ctcacatcgt tggatacccc cgcattgggtc ccaaggagag ctcaagttcg 180  
ctctcgagtc tttctgggat ggcaagagca gcgccgagga tttgcagaag gtggctgctg 240  
atctcaggtc atccatctgg aagcagatgg ctggtgctgg g 281

<210> 2774  
<211> 286  
<212> nucleic acid  
<213> Glycine max  
<400> 2774



gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctgcttca acttgctccc 60  
tcagaagcga agaagaagcc acagagaact agtctcctac tctcaccgcg aagaaaaaaa 120  
tggcatctca catcgttgga taccctcgta tgggtcccaa gagagagnnc aagttcgctc 180  
tcgagtcttt ctgggatggc aagagcagcg ccgaggattt gcagaagggtg tcttctgac 240  
tcagggcacc atctggaagc agatggctga tgctgggac aagtac 286

<210> 2775  
<211> 310  
<212> nucleic acid  
<213> Glycine max

<400> 2775

cncngcacnc gtacgtaagc tcggaattcg gctcgagctt gtcctctcag angcgaagaa 60  
gaagccacag agaantagtc tctactctc acccgccaag anaaaaatgg cattctcaca 120  
tcogttggat accctcgat ggggtcccaag agagagctca agttcgctct cgagtcttct 180  
gggatggcaa gnnccagcgc gaggatttgc agaagggtgc tctgatctca gggcatccat 240  
ctggaagcag atggctgatg ctgggatcaa gtacatcccc agcaacattt ctctcatatg 300  
accaggttct 310

<210> 2776  
<211> 299  
<212> nucleic acid  
<213> Glycine max

<400> 2776

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tcagaancga agaagaagcn acagagaact agtctcctac tctccaccgc caagaaaaaa 120  
atggcatctc acatcgttgg ataccctcgt atgggtccca agagagagct acaagttcgc 180  
tctcgagtct ttctgggatg gcaagagcag cgcgaggat ttgcagaagg tgtcttctga 240  
nctnagggca tccatctgga agcagatggc nnatgctggg atcattacat cccagcaa 299

<210> 2777  
<211> 253  
<212> nucleic acid  
<213> Glycine max

<400> 2777

cttgctcnnt naganncgaa gaagaagcca cagaggacta gtctnctac tgctcaccgc 60  
caagaaaaaa atggcatctc anacccgttg gatacccccg tatgggtcnc aanagagagc 120  
tcaagttngc tgctcgagtc tttctgggat ggcaatagca gngccganga tttgcagaag 180  
gtgtcttctg atctcagggc atccatctgg aatcagatgg ctgatgcngg gatcaagtnc 240  
atccccagca aca 253

<210> 2778

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 2778

gcaaangcac gcgagacgta anctcggaat tcggctcgag naacttgntc cctcagnagc 60  
gaagaagaag cnacananaa ctagtctcct acnncnaacc gnaagaaaaa natggcatct 120  
cacatcggtg gatacccccg tatnggtngc aanagagagc tcaagttcgc tctcgagtct 180  
tnctgngatg gnaagannag cgccgaggat ttgcagaagg tgtcttctga tctcagggca 240  
tccatctgga agcagatggc tangctggga tcaagtacat cccangca 288

<210> 2779

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2779

aaatnctana agtcgcangc acgcgnacgt aanctcgga ntcggctcga gnaacttgct 60  
ccctcagaag cgaagaagaa gccacanaga actagtctcc tactgctcac ccgcnagaaa 120  
aaaatggcat ctncacatnc gttggatanc cccgnatgng ngcccaagan agagctcaag 180  
ttcgctctcg agtctttctg ggatggcagn agcagcgccg aggatttgca gaaggtgtct 240  
tcngantca gggcatccat ctngaagcag atngctgatg ctgggatcaa gnacatctcc 300  
aggaacactt tctctnactn 320

<210> 2780

<211> 249

<212> nucleic acid

<213> Glycine max

<400> 2780

ctgcntcaac ttgctccctg caganncgaa ncaagaagcc acagagnact agtctnccta 60  
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aaganagagn tcaagttcgc tctcgagtct ttctgggatg gnaagagcag cgccgaggat 180  
ttgcagaagg tgtnttctga tctcagggna tncatctgga agnagatggc tgatgntggg 240  
ntcaagtac 249

<210> 2781

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 2781

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gaagaagcca cagagaacta gtctcctact notcaccgc aagaaaaaaa tggcatctca 120  
catacgttgg atacccccgn atgggtccca agagagagct caagtctcgt ctcgagtctt 180  
tctgggatgg caagagcagc gccgaggatt tgcagaaggt gtctcngatc tcagggcatc 240  
catctggaag cagatggctg atgctgggat nagtacann ccagcaacat ttctctcata 300

<210> 2782

<211> 262

<212> nucleic acid

<213> Glycine max

<400> 2782

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gggcccaaga gagagctcaa gttcgctctc gantctttct gggatggcaa gagcagcgcc 180  
gaggatttgc agaaggtgtc tctgatctca gggcatccat ctggaagcag atgctgatgt 240  
ggtcaagnac tcccgcacan tt 262

<210> 2783

<211> 242

<212> nucleic acid

<213> Glycine max

<400> 2783

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aagaagaagc cacagagaac cagtctccta ctctctctac acccacaaga aaaatggcat 120  
ctcacatcgt tggatacccc cgcattgggtc ccaagagaga gctcaagttc gctctcgagt 180  
ctttctggga tggcaagagc agcgccgagg atttgcagaa ggtggctgct gatctcaggt 240  
ca 242

<210> 2784

<211> 308

<212> nucleic acid

<213> Glycine max

<400> 2784

ncgcntgcac gcgtacgtaa gtcggaatt cggctcgaga acttgctccc tcagaagcga 60  
agaagaagcc acagagaact agtctctctac tcttcacccg caagaaaaaa atggcatctc 120  
acatcgttgg atacccccgt atgggtccca agagagagct caagtctcgt ctcgagtctt 180  
tctgggatgg caagagcagc gccgaggatt tgcagaaggt gtcttctgat ctcagggcat 240  
ccatctgnan canatggctg atnctgggnt ncagtacatc cccagcaaca tttcttctca 300  
tatgacca 308

<210> 2785

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2785

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ctcaagttcg ctctcgagtc tttctgggat ggcaagagca gngccgagga tttgcagaag 180  
gtgtctctga tctcagggca tccatctgg nagcagatgg ctgatgtggg atnngtacat 240  
cccagcaaca tttctcncat cngacangtt ctcgacg 277

<210> 2786

<211> 280  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2786  
  
 tatnaacttg ctcnntcaga ncgcgaaagaa gaagnacacag agaanchagt ctccctantnc 60  
 tcaccccgnaa gnnaaaaatg ggcattctcn catcggttga taacccccgt atgggtncca 120  
 agagagagct caagttcgnt ctncagtcct tctgggatgg caagagcagc gncngaggat 180  
 ttgcagaagg tgtcttctga tctcaggnc tccatctgga agcagatggc tgntgntggg 240  
 atcagtacat cccnagcaac acttctctca ctatgaccag 280

<210> 2787  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2787  
  
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 ccacananaa ccagtctcct actctctctc acccacaaga naaatgggat ctcacatcgt 120  
 tggntacccc cgcattgggtt cccaaganag agctcaagtt cgctctcgag tctttctggg 180  
 ntggcnagag cagcgccgag gatttgcaga aggtggctgc tgatctcagg tcatccaatc 240  
 tggaancaag attgccngat cggggatcaa gctccatncc cagcaacann tttttgct 298

<210> 2788  
 <211> 151  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2788  
  
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 gctgggatca agtacatccc cagcaacact ttctctcact atgaccaggt tctcgacgcc 120  
 accgccaccc tcggtgccgt tccaccaagg t 151

<210> 2789  
 <211> 234  
 <212> nucleic acid  
 <213> Glycine max

<400> 2789

gcacgcgtac gtaagctcgg aattcggctc gagcttcaac ttgctccctc agaagcgaag 60  
aagaagccac agagnactag tctctactct tcaccogcaa naaaaaaaaatg ggcatctcac 120  
attcgttgga tcccccgta tgggtcccaa gagagagctc aagttcgctc tcgagtcttc 180  
naggatggca agagcagcgc cgaggatttg cagaaggtgt cttctgatct cagg 234

<210> 2790

<211> 138

<212> nucleic acid

<213> Glycine max

<400> 2790

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cacatcnttg gatacccccg tatgggtccc aagagagagc tcaagttcgc tctcgagtct 120  
ttctgggatg gcaagagt 138

<210> 2791

<211> 152

<212> nucleic acid

<213> Glycine max

<400> 2791

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atcaagtaca tccccagcaa cactttctnn tnactatgac caggttctcg acgccacccg 120  
ccaccctcgg tgccgttcca cncangnnag gn 152

<210> 2792

<211> 501

<212> nucleic acid

<213> Glycine max

<400> 2792

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tcggaattcg gctcgagcaa cnttccaatc cttccaacca ccactantgg atccttonct 120  
canacngtag aactgaggag ggtacgcctg gagttcaagg ctaacatgat ctccgaggaa 180  
gagtatgtta agtcaattaa ggaggaaatt cncaaagttg ttgaacttca anaagagctt 240

gatattgatg ttcttgttca tggagaacca gagagaaatg atatggttga gtacntcggt 300  
gagcaattgt caggctttgc cttcactgtt aatgggtggg tgcaatccta tggttcccgt 360  
tgtgtgaagc cacnaatcat ctatggtgat gtgagccgcc caaagccaat gactgtcttc 420  
tggtcancctc tggctcagan ctttaccag cgccnaatga agggaatgct taccggtccg 480  
gttaccaatc ccaactgggn c 501

<210> 2793  
<211> 412  
<212> nucleic acid  
<213> Glycine max  
<400> 2793

gagaagacga cagaaggggg ctaaggcatt gtctggcaac aaggatgtgg ctttcttctc 60  
tgctaattgt gcanctcang cttcaaggaa gtccctctcca anagtgacca acnaggctgt 120  
tcagaaggct gctgctgcat tgaagggttc agatcatcnc cgtgcaacaa atgtcagtgc 180  
cagactggat gctcaacaaa anaagctcaa ccttccaatc cttccaacca ccactattgg 240  
atccttccct cagactgtan aactgaggag ggtacgcogt gagttcaagg ctaacaagat 300  
ctccgaggaa gagtatgtta agtcaattaa ggaggaaatt cgcaaagttg ttgaacttca 360  
agaagagctt gatattgatg ttcttgttca tgganaacca anaganaaat ta 412

<210> 2794  
<211> 350  
<212> nucleic acid  
<213> Glycine max  
<400> 2794

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gcatttgctg cacaaaaaat tgttgaagtt aacgcattgg ctaaggcatt gtctggcaac 120  
aaggatgtgg ctttcttctc tgctaattgt gcagctcagg cttcaaggaa gtccctctcca 180  
agagtgacca acgaggctgt tcagaaggct gctgctgcat tgaagggttc agatcatcgn 240  
cgtgcaacaa atgtcagtgc cagactggat gctcaacaaa agaagctcaa ccttccatcc 300  
ttccaaccac cactattgga tccttccctc agactgtaga actgaggagg 350

<210> 2795

<211> 454  
 <212> nucleic acid  
 <213> Glycine max

<400> 2795

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agaagctcaa ngttccngtc cttccaacca ccactattgg atccttccct cagactgtag 120
aactgangag ggtacgccgt gagttcaagg ctaacaagat ctccgaggaa gagtatgtta 180
agtcaattaa ggaggaaatt cgcaaagttg ttgaacttca anaagagctt gatattgatg 240
ttcttgttca tggagaacca gagagaaatg atntgggtga gtacttcggt gagcaattgt 300
caggctttgc cttcactggt aatgggtggg tgcaatctat ggggccggt gtgtgaaagc 360
caccaattca tctatgggtg aatgtgnaaa nncngtccaa aagccaatga ctgtcttctg 420
gtcatctntg gcttaaangc tttaccaaag cgct 454
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<210> 2796  
 <211> 446  
 <212> nucleic acid  
 <213> Glycine max

<400> 2796

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tgtcagtgcc agactggatt ctcaacaaaa gaagctcaac cttccaatcc tgccaaccac 120
cactattgga tcttccctc agactgtaga actgaggagg gtacgccgtg aattcaaggc 180
taacaagatc tccgaggaag agtatgtaaa gtcaattaag gaggaaattc gcaaagttgt 240
tgagcttcaa gaagagcttg atattgatgt tcttgggtcat ggagaaccag agagaaatga 300
tatgggtgag tactttcggg gaacaattgt caagctttgg cntnaccggt aatgggtngg 360
tgcaatccta tggttcccggt tcgtgaaanc cccgatcatt tatgggaatg ttagccgccc 420
aaagccatga ccgntttttg gtattt 446
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<210> 2797  
 <211> 489  
 <212> nucleic acid  
 <213> Glycine max

<400> 2797



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 ctgagaccac ctaccagatt gctttgtcta tcaaggacga agtggaagac cttgaaaagg 120  
 ctggcatcac tggtatccaa attgatgaag ctgctttgag agaggggtctt ccaactgagga 180  
 aatcagagca agctcactac ttggactggg ctgtccatgc cttcagaatc accaatgttg 240  
 gtgtccagga taccacccag gtacactctt ttggatcatc gcaaatact gaattagaaa 300  
 ttttttttgt tcactctcat tttcacatat gttgtaataa tcaacttttc gtattgacag 360  
 atccacactc acatgtgcta ctcgaaactt aacgacatca tccactccat catcgacatg 420  
 gacgccgatg ttatcaccat tgagaaaatc cgccccgacg anaancttcc gtcagtcctc 480  
 cgccaangg 489

<210> 2798  
 <211> 340  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2798

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 tgctgccccaa aaaattgttg aagttaacgc attggctaaa gcattgtctg gccacaagga 120  
 tgangccttc ttctctggta atgctgctgc tctggcttca aggaagtctt ctccaagagt 180  
 gaccaacgag gctgttcaga aggctgctgc tgcattgaag ggttcagatc atcgccgtgc 240  
 aanaaatgtc agtgccagac tggattctca acaaaagaag ctcaaccttc caatcctgcc 300  
 aaccaccact attggatcct tccctcagac tgtagaactg 340

<210> 2799  
 <211> 317  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2799

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 tcggtgagca attgtcaggc tttgccttca ctgttaatgg gtgggtgcaa tcctatgggt 120  
 cccgttgtgt gaagccacca atcatctatg gtgatgtgag ccgccccaaag ccaatgactg 180  
 tcttctggtc atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg 240

gtcctgttac cattctcaac tggtcctttg ttagaaatga ccaacctaga tctgagacca 300  
cctaccagat tgctttg 317

<210> 2800  
<211> 317  
<212> nucleic acid  
<213> Glycine max  
<400> 2800

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ggaaatcaga acaagctcac tacttggact gggctgtcca tgccttcaga atcaccaatg 180  
ttgggtgtgca ggataccact cagatccaca cccacatgtg ctactccaac ttcaacgaca 240  
tcatccactc catcatcgac atggacgtg atgttatcac cattgagaac tctcgctccg 300  
atgagaagct cctgtca 317

<210> 2801  
<211> 337  
<212> nucleic acid  
<213> Glycine max  
<400> 2801

cnngtcgan gcacgcntac gtaagctcgg aattcggctc gaggccacca atcatctatg 60  
gtgatgtgag ccgccc aaag ccaatgactg tcttctggtc atctctggct cagagcttta 120  
ccaagcnccc aatgaaggga atgcttaccg gtctgttac cattctcaac tggtcctttg 180  
ttagaaatga ccaacctaga tctgagacca cctaccagat tgctttggct atcaaggacg 240  
aagtggagga ccttgaaaag gctggcatca ctgttatcca aattgatgaa gctgctttga 300  
gagaggggtct gccactgagg aaatcagaac aagctcn 337

<210> 2802  
<211> 329  
<212> nucleic acid  
<213> Glycine max  
<400> 2802

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atcgccgtgc aacaaatgtc agtgccagac tggatgctca acaaaagaag ctcaaccttc 120  
 caatccttcc aaccaccact attggatcct tccctcagac tgtagaactg aggagggtag 180  
 gccgtgagtt caaggctaac aagatctccg aggaagagta tgттаagtca attaaggagg 240  
 aaattcgcaa agttgttgaa cttcaagaag agcttgatat tgatgttctt gttcatggag 300  
 aaccagagag aatgatatg gttgagtac 329

<210> 2803  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2803

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 cactacttgg actgggctgt ccatgccttc agaatcacca atgttggtgt ccaggatacc 120  
 acccagatcc aactcacat gtgctaactg aacttcaacg acatcatcca ctccatcatc 180  
 gacatggacg ccgatgttat caccattgag aactctcgct ccgacgagaa gcttctgtca 240  
 gtcttccgcg aaggtgtgaa gtatggtgct ggaattggcc ctggtgtcta tgacatccac 300  
 tccccaaaga tacc 314

<210> 2804  
 <211> 328  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2804

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 aacaagatct ccgaggaaga gtatgtaaag tcaattaagg aggaaattcg caaagttggt 120  
 gagcttcaag aagagcttga tattgatgtt cttgttcatg gagaaccaga gagaaatgat 180  
 atggttgagt acttcggtga acaattgtca ggctttgcct tcaccgttaa tgggtgggtg 240  
 caatcctatg gttcccggtg cgtgaagcca ccgatcatct atggtgatgt gagccgcca 300  
 aagccaatga ccgtcttctg gtcattctc 328

<210> 2805  
 <211> 323  
 <212> nucleic acid

<213> Glycine max

<400> 2805

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gatgtggcct tcttctctgc taatgctgca gctcaggcctt caaggaagtc ctctccaaga 120  
gtgaccaacg aggcgtgtca gaaggctgct gctgcattga agggttcaga tcatcgccgt 180  
gcaacaaatg tcagtgccag actggatgct caacaaaaga agctcaacct tccaatcctt 240  
ccaaccacca ctattggatc cttccctcag actntagaac tgaggagggt acgccgtgag 300  
ttcaaggcta acaagatctc cga 323

<210> 2806

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2806

ngnncgcatg cacgcgtacg tnaagctcgga attcggctcg agatatgggtt gagtacttcg 60  
gtgagcaatt gtnaggcttt gccttcaactg ttaatgggtg ggtgcnatcc tatggttccc 120  
gttgtgtgaa gccaccaatc atctatgggtg atgtgagccg cccaaagcca atgactgtct 180  
tctggtcacg tctggctcag agctttacca agcgcccaat gaagggaatg cttaccggtc 240  
ctgttaccat tctcaactgg tcctttgtta ganatgacca acctagatct gagaccacct 300  
accagattgc tt 312

<210> 2807

<211> 295

<212> nucleic acid

<213> Glycine max

<400> 2807

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gtgagcaatt gtcaggcttt gccttcaactg ttaatgggtg ggtgcaatcc tatggttccc 120  
gttgtgtgaa gccaccaatc atctatgggtg atgtgagccg cccaaagcca atgactgtct 180  
tctggtcacg tctggctcag agctttacca agcgcccaat gaagggaatg cttaccggtc 240  
ctgttaccat tctcaactgg tcctttgtta gaaatgacca acctagatct gagac 295



gaaatgatat ggttgagtac ttcggtgagc aattgtcagg ctttgccctc actgttaatg 240  
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 gccgcc 306

<210> 2811  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2811

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 tgaaggtgtg aagtatggtg ctggaattgg ccctggtgtc tatgacatcc actccccaag 180  
 aataccacca actgaagaaa tcgctgacag aatcaataag atgcttgacg tgctcgagaa 240  
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 gccagccctc 310

<210> 2812  
 <211> 353  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2812

nnnnnnaaaa gtcgggaagt cgcangcacg cgtacgtaag ctcggaattc ggctcgnanc 60  
 tcgagccgaa tcggctcgag ccagattgct ttggctatca aggaagnagt ggaggacctt 120  
 acnaaggctg gcatcactgt actccaaatt gatgaagctg ctttgagaga gggctctgcca 180  
 ctgaggaaat cagaacaagc tcactacttg gactgggctg tccatgcctt cagaatcacc 240  
 aatgttggtg tgcaggatac cactcagatc cacaccacaca tgtgctactc caacttcaac 300  
 gacatcatcc actccatcat cgacatggac gctgatgtta tcaccattga gat 353

<210> 2813  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2813

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actgaggaaa tcagaacaag ctactactt ggactgggct gtccatgcct tcagaatcac 120  
caatgttggt gtgcaggata ccactcagat ccacaccac atgtgctact ccaacttcaa 180  
cgacatcatc cactccatca tcgacatgga cgctgatgtt atcaccattg agaactctcg 240  
ctccgatgag aagctcctgt cagtcttccg tgaagggtgt aagtatgggt ctggaat 297

<210> 2814  
<211> 551  
<212> nucleic acid  
<213> Glycine max  
<400> 2814

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agaactctcg ctccgatgag aagctcctgg tcagtcttcc gtgaagggtgt gaagtatgggt 180  
gctggaattg gccctgggtgt ctatgacatc cactccccaa gaataccacc aactgaagaa 240  
atcgctgaca gaatcaataa gatgcttgca gtgctcgaga agaacatctt gtgggtcaac 300  
cctgactgtg gtctcaagac ccgcaagtac actgaagtga agccagccct cacaacatg 360  
gttgccgcag caaaactcat ccgtaacgaa cttgccaagt gaatgggtata aagaaagtag 420  
aatctacaag ttcatgtggt ctgcttttat tataccncca aggaaaaatt ttctatantn 480  
gggtggttca aataaccggt gtggaatatt tanaggttta acatgctctg tgagcaattg 540  
atctttctca c 551

<210> 2815  
<211> 336  
<212> nucleic acid  
<213> Glycine max  
<400> 2815

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tccaatcctg ccaaccacca ctattggatc cttccctcag actgtagaac tgaggagggt 180  
acgccgtgaa ttcaaggcta acaagatctc cgaggaagag tatgtaaagt caattaagga 240

ggaaattcgc aaagttgttg agcttcaaga agagcttgat attgatgttc ttgttcacgg 300  
agaaccagag agaaatgata tggttgagta cttcgg 336

<210> 2816  
<211> 313  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2816

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catccactcc atcatcgaca tggacgccga tgttatcacc attgagaact ctcgctccga 120  
cgagaagctt ctgtcagtct tccgcgaagg tgtgaagtat ggtgctggaa ttggccctgg 180  
tgtctatgac atccactccc caagaatacc accaactgaa gaaattgctg acagaatcaa 240  
caagatgctg gcagtgctcg agangaacat cttgtgggnt gaacctgact gtgggctcaa 300  
gaccgcgtaaa gtn 313

<210> 2817  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2817

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ttccactgag gaaatcagag caagctcact acttggactg ggctgtccat gccttcagaa 120  
tcaccaatgt tgggtgtccag gataccaccc agatccacac tcacatgtgc tactcgaact 180  
tcaacgacat catccactcc atcatcgaca tggacgccga tgttatcacc attgagaact 240  
ctcgctccga cgagaagctt ctgtcagtct tccgcgaagg tgtgaagtat ggtgctggaa 300  
ttgg 304

<210> 2818  
<211> 438  
<212> nucleic acid  
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 atcaaggacn aantggaaga ccttgaaaag ggctggcatc actgntatcc aaattgatga 360  
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 gaccaagttg gatgatgaga tcaagtcatg gctagctttt gctgccc aaaattgttga 180  
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 ctgtcagtct tccgcgaagg tgtgaagtat ggtgctggaa ttggccctgg tgtctatgac 240  
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 acatcatcca ctccatcatc gacatggacg ctgatgttat caccattgag aactctcgct 180  
 ccgatgagaa gtcctgtca gtcttccgtg aagggtgtgaa gtatgggtgct ggaattggcc 240  
 ctggtgtcta gacatccact cccaagaat accaccaact gaagaaatcg 290

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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2823  
  
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 atgtggcctt cttctctgct aatgctgcag ctcaggcttc aaggaagtcc tctccaagag 180  
 tgaccaacga ggctgttcag aaggctgctg ctgcattgaa gggttcagat catcgccgtg 240

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265

<210> 2824  
<211> 289  
<212> nucleic acid  
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cgtcgcatgc acgcgtacgt aagctcggaa ttcggctcga gcaaagttgt tgaacttcaa 60  
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tacttcggtg agcaattgtc aggccttggc ttcactgtta atgggtgggt gcaatcctat 180  
ggttcccggt gtgtgaagcc accaatcatc tatggtgatg tgagccgccc aaagccaatg 240  
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CGTCG  
CATGC  
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TACGT  
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GCAAG  
TTGT  
TGAAC  
TTCAA  
GAAGA  
GCTTG  
ATATT  
GATGT  
TCTTG  
TTCAT  
GGAGA  
ACCAG  
AGAGA  
AATGA  
TATGG  
TTGAG  
TACTT  
CGGTG  
AGCAA  
TTGTC  
AGGCT  
TTGGC  
TTCAC  
TGTTA  
ATGGG  
TGGGT  
GCAAT  
CCTAT  
GGTT  
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<210> 2825  
<211> 265  
<212> nucleic acid  
<213> Glycine max  
  
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gctgcagctc aggccttcaag gaagtcctct ccaagagtga ccaacgaggc tgttcagaag 120  
gctgctgctg cattgaaggg ttcagatcat cgccgtgcaa caaatgtcag tgccagactg 180  
gatgctcaac aaaagaagct caaccttcca atccttccaa ccaccactat tggntccttc 240  
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<210> 2826  
<211> 304  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2826

agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gcttcacact gctgttgatc 60  
nagttaacga gaccaagttg gatgatgaga tcaagtcatg gctagctttt gctgccccaa 120  
aaattgttga agttaacgca ttggctaaag cattgtctgg ccacaaggat gaggccttct 180  
tctctggtaa tgctgctgct ctggcttcaa ggaagtcttc tccaagagtg accaacgagg 240

ctgttcagaa ggctgctgct gcattgaagg gttcagatca tcgccgtgca acaaattgtca 300  
gtgc 304

<210> 2827  
<211> 269  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2827

ctggtttgan tgttcttggt gagacctact ttgctgacat ccctgctgag gcatacaaga 60  
ccctcacatc tctgaatggc gtcactgcat atggatttga tttggtccgt ggaaccaaca 120  
ctcttgattt gatcaagggg ggatttccca gcggaaaata cctctttgct ggagtgggtg 180  
atggaaggna catctggggc aatgaccttg ctgcttctct cactaccttg cagggctctg 240  
agggcattgt gggcaaagat aagcttggt 269

<210> 2828  
<211> 315  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2828

tncatnnag cgnaggtann tacgtaagct cggaattcgg ctcgagtacg gctgcgagaa 60  
gacgacagaa gggcatggag aaccagagag caatgatatg gttgagtact tcggtgagca 120  
attgtcaggc ttgcttca ctgttaatgg gtgggtgcaa tcctatgggt cccgttgtgt 180  
gaagccacca atcatctatg gtgatgtgag ccgcccagg ccaatgactg tcttctggtc 240  
atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg gtctgttac 300  
cattctcaac tggtc 315

<210> 2829  
<211> 320  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2829

gtcgcangca cgcgtacgtn agctcggnat tcggtcgag caagcgccca atgaaggga 60  
tgcttaccgg tctgttacc attctcaact ggtcctttgt tagaaatgac caacctagat 120

ctgagaccac ctaccagatt gctttgtcta tcaaggacga ngtggaagac cttgaaaagg 180  
 cggcatact gttatccaaa ttgatgaagc tgctttgaga gaggtcttc cactgaggaa 240  
 atcagagcaa gctcactact tggactgggc tgtccatgcc ttcagaatca ccaatgttgg 300  
 tgtccaggat accaccaga 320

<210> 2830  
 <211> 512  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2830

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 gctgacctta aggcagctgg tgcttcatgg attcagtttg atgagcctac ccttgtcttg 180  
 gaccttgagt ctcaaaagt gcaagcattc actgacgcat atgcagaact tgcgcctgct 240  
 ttgtctgggt tgaatgttct tgttgagacc tactttgctg acatccctgc tgaggcatac 300  
 aagaaccctt acatnttctt gaatggcgtc actgcatatg gatttgattt ggtccgtgga 360  
 accaacactc ttgatttgat caaggttgga tttccagcgg aaaatacctc tttcttgga 420  
 tgggttgatg gaaggacatt tgggccaatg accttgctgg tttttttaat acctgcaggg 480  
 cttgaagggc atttgggcca aaataacctg tt 512

<210> 2831  
 <211> 325  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2831

nnngtcgcan gcacgcgtac gtaagctcgg aattcggctc gaggttgagt acttcggtga 60  
 acaattgtca ggctttgcct tcaccgttaa tgggtgggtg caatcctatg gttcccgttg 120  
 cgtgaagcca ccgatcatct atggtgatgt gagccgcca aagccaatga ccgtcttctg 180  
 gtcatctctg gctcagagct ttaccaagcg cccaatgaag ggaatgotta ccggtcctgt 240  
 taccattctc aactggctct ttgttagaaa tgaccaacct agatngagac cactaccaga 300  
 ttgctttgtc tatcaaggac gaagg 325

<210> 2832  
 <211> 323  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2832  
  
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 tntcgccgt gcaacaaatg tcagtgccag actggatgct caacaaaaga agctcaacct 180  
 tccaatcctt ccaaccacca ctattggntc cttccctcag actgtagaac tgaggaggga 240  
 ccgccgtgag ttcaaggcta acaagatctc cgaggaagag tatgttaagt caattaagga 300  
 ggaaattcgc aaagttgttg aan 323

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<210> 2833  
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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2833  
  
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 cgcaaagttg ttgaacttca agaagagctt gatattgatg ttcttggttca tggagaacca 120  
 gagagaaatg atatggttga gtacttcggt gagcaattgt caggctttgc cttcactggt 180  
 aatgggtggg tgcaatccta tggttcccgt tgtgtgaagc caccaatcat ctatggtgat 240  
 gtgagccgcc caaagccaat gactgtcttc tggctcatctc tggctcagag ctttac 296

<210> 2834  
 <211> 265  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2834  
  
 cgcccaatga agggaatgct taccggtcct gttaccattc tcaactggtc ctttggttaga 60  
 aatgaccaac ctatgcttga gaccacctac cagattgctt tgtctatcaa ggacgaagtg 120  
 gaagaccttg aaaaggctgg catcactggt atccaaattg atgaagctgc tttgagagag 180  
 ggtcttccac tgaggaaatc agagcaagct cactacttgg actgggctgt ccatgccttc 240

agaatcacca atgttggtgt ccagg

265

<210> 2835  
<211> 325  
<212> nucleic acid  
<213> Glycine max

<400> 2835

wgtngcnngc acgcgtacgt aagctcggaa ttccggctcga gttcccagtg gaaaataacct 60

ctttgctgga gtggttgatg gaaggaacat ctggggccaat gaccttgctg cttctctcac 120

tacattgcag ggtcttgagg gcattgtggg caaagataag cttgttggtt ccacctctc 180

ctcccttctt cacactgctg ttgatcttgt taacgagacc aagttggatg acgagatcaa 240

gtcatggcta gcatttgctg cacaaaaaat tgttgaagtt aacgcattgg ctaaggcatt 300

gtctggcaac aaggagtggc cttct 325

<210> 2836  
<211> 346  
<212> nucleic acid  
<213> Glycine max

<400> 2836

gttaacgcat tggctaaggc attgtctggc aacaaggatg tggccttctt ctctgctaatt 60

gctgcagctc aggcttcaag naagtcctct ccnagagtga ccaacgaggc tgttcagaag 120

gctgctgctg cattgaaggg ttcanatcat cgccgtgcaa caaatgtcag tgccagactg 180

gatgctcaac aaaagaagct caaccttcca atccttccaa ccaccactnt tgnntccttc 240

cctcagactg tagaactgag gagggtagc gtgagttcaa ggtaacaaga ntccgaggaa 300

gagtatgtta agnccattaa ggaganatnt caagtgtgaa ctnaag 346

<210> 2837  
<211> 312  
<212> nucleic acid  
<213> Glycine max

<400> 2837

tcgcangcac gcgtacgtaa gctcgggaatt cggctcgagg accttgctgc ttctctcact 60

acattgcagg gtcttgaggg cattgtgggc aaagataagc ttgttggtgc cacctcctcc 120

tcccttcttc acactgctgt tgatcttggt aacgagacca agttggatga cgagatcaag 180  
 tcatggctag catttgctgc acaaaaaatt gttgaagtta acgcattggc taaggcattg 240  
 tctggcaaca aggatgtggc cttcttctct gctaattgctg cagctcaggc ttcaaggaag 300  
 tcctctccaa ga 312

<210> 2838  
 <211> 307  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2838

ntatccntnn acgtcgcang cacgcgtacg taagctcgga attcggctcg agngagggta 60  
 cgccgtgagt tcaaggctaa caagatctcc gaggaagagt atgttaagtc aattaaggag 120  
 gaaattcgca aagttgttga acttcaagaa gagcttgata ttgatgttct tgttcatgga 180  
 gaaccagaga gaaatgatat ggttgagtac ttccgtgagc aattgtcagg ctttgccttc 240  
 actgttaatg ggtgggtgca atcctatggt tcccgttggt tgaagccacc aatcatctat 300  
 ggtgatg 307

<210> 2839  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2839

cttnnagtcn catncacnag tacgnaantc ngcnnnncng ttcttggtga gacctacttt 60  
 gctgacatcc ctgctgaggc atacaagacc ctacatctc tgaatggcgt cactgcatat 120  
 ggatttgatt tgggtccgtgg aaccaacact cttgatttga tcaaggggtg atttcccagc 180  
 ggaaaatacc tctttgctgg agtggttgat ggaaggaaca tctgggcca tgaccttgct 240  
 gcttctctca ctaccttgca gggctttgag ggcattgtgg gcaaagataa gcttgttgtg 300  
 tccacctcct 310

<210> 2840  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max



<400> 2840

nttcgcatgc acgcgtacgt aagctcggaa ttccggctcga gcacctacca gattgctttg 60

tctatcaagg acgangtgga agaccttgaa aaggctggca tcaactgttat ccaaattgat 120

gaagctgctt tgagagaggg tcttccactg aggaaatcag agcaagctca ctacttggac 180

tgggctgtcc atgccttcag aatcaccaat gttgggtgtcc aggataccac ccagatccac 240

actcacatgt gctactcgaa cttcaacgac atcatccact ccatcatcga catggac 297

<210> 2841

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2841

gtcgtttnat gcaacgcgtac gtaagctcgg aattcggctc gagaaaaatt gttgaagtta 60

acgcattggc taaggcattg tctggcaaca aggatgtggc cttcttctct gctaattgctg 120

cagctcaggc ttcaaggaag tcctctccaa gagtgaccaa cgaggctgtt cagaaggctg 180

ctgctgcatt gaagggttca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatg 240

ctcaacaaaa gaagctcaac cttccaatcc ttccaaccac cactattgga tccttccctc 300

aga 303

<210> 2842

<211> 241

<212> nucleic acid

<213> Glycine max

<400> 2842

gttcttgttc atggagaacc agagagaaat gatatggttg agtacttcgg tgaacaattg 60

tcaggctttg cttcacctgt taatgggtgg gtgcaatcct atggttcccg ttgcgtgaag 120

ccaccgatca tctatggtga tgtgagccgc ccaaagccaa tgaccgtctt ctggtcattc 180

ctggctcaga gctttaccaa gcgcccaatg aagggaatgc ttaccgggtc tgttaccatt 240

c 241

<210> 2843

<211> 296

<212> nucleic acid

<213> Glycine max

<400> 2843

ntgcngcgta ngtaagctcg gaattcggct cgagtcaggc cttctttctct gctaagtctg 60  
 cagctcaggc ttcaaggaag tcctctccaa gagtgaccaa cgaggctggt cagaaggctg 120  
 ctgctgcatt gaagggttca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatg 180  
 ctcaacaaaa gaagctcaac cttccaatcc ttccaaccac cactattggn tccttccctc 240  
 agactgtaga actgaggagg gtacgccgtg agttcaaggc tancaagatc tccgag 296

<210> 2844

<211> 265

<212> nucleic acid

<213> Glycine max

<400> 2844

cgcccaatga agggaaatgct taccggctcct gttaccattc tcaactggtc ctttgttaga 60  
 aatgaccaac ctagatctga gaccacctac cagattgntt tgtctatcaa ggacgaagtg 120  
 gaagaccttg aaaaggctgg catcactgtt atccaaattg atgaagctgc tttgagagag 180  
 ggtcttccac tgaggaaatc agagcaagct cactacttgg actgggctgt ccatgccttc 240  
 agaatcacca angttggtgt ccagg 265

<210> 2845

<211> 315

<212> nucleic acid

<213> Glycine max

<400> 2845

atntgtcgca tgcncgcgta cgtaagctcg gnattcggct cgagngacat catccactcc 60  
 atcatcgaca ggncgccgat gttatcacca ttgagaantc tcgctccgac gagangcttc 120  
 tgtcagtctt ccgcgaaggt gtgangtatg gtgctggaat tggccttgggt gtctatgaca 180  
 tccactcccc aagaatacca ccaactgaag anattgctga cagaatcaac aagatgctgg 240  
 cagtgtctga gaagnacatc ttgtgggtga nccctgactg tgggctcaag acccgtaagt 300  
 aactgaggt gaagc 315

<210> 2846

<211> 311  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2846  
  
 ngncatgcac gcgtacgtaa gctcgggaatt nggctcgagt cagatcatcg ccgtgcaaca 60  
 aatgtcagtg ccagactgga ttctcaacaa aagaagctca accttccaat cctgccaaacc 120  
 accactattg gatccttccc tcagactgta gaactgagga gggtaacgacg tgaattcaag 180  
 gctaacaaga tctccgagga agagtatgta aagtcaatta angaggaant tcgcaaagtt 240  
 gttgagcttc aagaagagct tgatatngat gttcttggtc atggagaacc agagagaaat 300  
 gatatggttg n 311

<210> 2847  
 <211> 256  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2847  
  
 gaagccaccg atcatctatg gtgatgtgag ccgcccacaaag ccaatgaccg tcttctggtc 60  
 atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg gtcctgttac 120  
 cattctcaac tggtcctttg ttgaaatga ccaacctaga tctgagacca cctaccagat 180  
 tgctttgtct atcaaggacg aagtggaaga ccttgaaaag gctggcatca ctgttatcca 240  
 aattgatgaa gctgct 256

<210> 2848  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2848  
  
 gtcgcatgca cgcgtacgta agctcggaat tcgggctcga ggatttgatc aagggtggat 60  
 ttcccagcgg aaaatacctc tttgctggag tggttgatgg aaggaacatc tgggccaatg 120  
 accttgctgc ttctctcact accttgcagg gtcttgaggg cattgtgggc aaagataagc 180  
 ttgttggtgc cacctcctcc tcccttcttc aactgctgt tgatctagtt aacgagacca 240  
 agttggatga tgagatcaag tcatggctag cttttgctgc caaaaaatt gttgaagtta 300

acgcatgg

308

<210> 2849  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2849

acaagnanna ngtacgtaag ctcggaatt cggtcgagc cttcttctct gctaattgctg 60  
cagctcaggc ttcaaggaag tcctctccaa gaggtagcaa cgaggctggt cagaaggctg 120  
ctgctgcatt gaaggggtca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatg 180  
ctcaacaaaa gaagctcaac cttccaatcc ttccaaccac cactattgga tccttccttc 240  
agactgtaga actgaggagg gtacgccgtg agttcaaggc taacaagatc tc 292

<210> 2850  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2850

cgccangcag cgtacgtaag ctcggaattc gggtcgagcc actgaggaaa tcgaacaagc 60  
tcactacttg gactgggctg tccatgcctt cagaatcacc aatgttggtg tgcaggatac 120  
cactcagatc cacaccacac tgtgtacttc caacttcaac gacatcatcc actccatcat 180  
cgacatggac gctgatgtta tcaccattga gaactctgcg tccgatgaga agctcctgtc 240  
agtcttccgt gaaggtgtga agtatgggtc tggaattggc cctgggtgtct at 292

<210> 2851  
<211> 327  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2851

gtcncangca cgcntacgta anctcggaat tcgggtcgag aaatgatatg gttgagtact 60  
tcgggtgaaca attgtcaggc ttgtccttca ccgttaatgg gtgggtgcaa tcctatgggt 120  
cccgttgctg gaagccaccg atcatctatg gtgatgtgag ccgccccaaag ccaatnaccg 180  
tcttctggtc atctctggct cagagcttta ccaagcgccc aatgaaggga atgcttaccg 240

gtcctgttac cattctcaac tggtcctttg ttagaaatga ccaacctang tataaactcc 300  
acaccgaaaa atgaacatca aggaggg 327

<210> 2852  
<211> 345  
<212> nucleic acid  
<213> Glycine max  
<400> 2852

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aatnggcgtc actgcatatg gggttgattt ggccgtgga acccatactc ttgatttgat 120  
caagggtgga tttcccagtg gaaaataact ctttgctgga gtgggtgatg gaaggnacat 180  
ctgggccaat gaccttgctg cttctctcac tacattgcag ggtcttgagg gcattgtggg 240  
caaagataag cttgttggtt ccacctctc ctcccttctt cacactgtg ttgatcttgt 300  
taacgagacc aagttggatg acgagatcaa gtcattggcta gcatt 345

<210> 2853  
<211> 309  
<212> nucleic acid  
<213> Glycine max  
<400> 2853

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tgcatgaag gggtcagatc atcgccgtgc acaaatgct agtgccagac tggatgctca 120  
acaaaagaag ctcaacctc caatccttcc aaccaccact attggatctt tccctcagac 180  
tgtagaactg aggagggtac gccgtgagtt caaggctaac aagatctccg aggaagagta 240  
tgttaagtca attaaggagg aaattcgcaa agttgttgaa cttcaagaag agcttgatat 300  
tgatgttct 309

<210> 2854  
<211> 311  
<212> nucleic acid  
<213> Glycine max  
<400> 2854

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 ttaccaagcg cccaatgaag ggaatgctta ccggtcctgt taccattctc aactgggtcct 180  
 ttgttagaaa tgaccaacct agatctgaga ccacctacca gattgctttg gctatcaagg 240  
 acgaatggag gaccttgaaa aggctggcat cactgttatc caaattgatg aagctgcttt 300  
 gagagagggg c 311

<210> 2855  
 <211> 324  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2855

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 cgctgatgtt atcaccattg agaactctcg ctccgatgag aagctcctgt ncagtcttcc 120  
 gtgaagggtg gaagtatggg gctggaattg gccctgggtg ctatgacatc cactccccaa 180  
 gaataccacc aactgaagaa atcgctgaca gaatcaataa gatgcttgca gtgctcgaga 240  
 agaacatctt gtgggtcaac cctgactgtg gttcaagacc cgcaagtaca ctgaagtga 300  
 gccagccctt nacaaacatg gttg 324

<210> 2856  
 <211> 311  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2856

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc cttgaaaagg ctgggcatca 60  
 ctgttatcca aattgatgaa gctgctttga gaganggtct tccactgagg aaatcagagc 120  
 aagctcacta cttggactgg gctgtccatg ccttcagaat caccaatgtt ggtgtccagg 180  
 ataccacca gatccacact cacatgtgct actcgaactt caacgacatc atccactoca 240  
 tcatcgacat ggacgccgat gtttcacat tgagaactct cgctccgacg agaagcttct 300  
 gtcagtcttc c 311

<210> 2857  
 <211> 258  
 <212> nucleic acid



<213> Glycine max

<400> 2860

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 cctgttacca ttctcaactg gtcctttgtt agaaatgacc aacctagatc tgagaccacc 120  
 taccagattg ctttgtctat caaggacgaa gtggaagacc ttgaaaaggc tggcatcact 180  
 gttatccaaa ttgatgaagc tgctttgaga gaggggtcttc cactgaggaa atcagagcaa 240  
 gctcactact tggactgggc tgtccatgcc ttcagaatca ccaatg 286

<210> 2861

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 2861

cgcatgcncg cgtacgtaag ctcggaattc ggctcgagaa gcattgtctg gncacaagga 60  
 tgaggccttc ttctctggta atgctgctgc tctggcttca aggaagtctt ctccaagagt 120  
 gaccaacgag gctgtncaga aggctgctgc tgcattgagg gttcagatca tcgccgtgca 180  
 acaaatgtca gtgccagact ggattctcaa caaaagaagc tcaaccttcc aatcctgcc 240  
 accaccacta ttggatcctt ccctcagact gtagaactga ggaggggtacg ccgtgaattc 300  
 aag 303

<210> 2862

<211> 311

<212> nucleic acid

<213> Glycine max

<400> 2862

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 ctccgatgag aagctcctgt cagtcttccg tgaagggtgtg aagtatggtg ctggaattgg 120  
 ccctgggtgc tatgacatcc actccccaag aataccacca actgaagaaa tcgctgacag 180  
 aatcaataag atgcnnngcag tgctcgagaa gaacatcttg tgggtcaacc ctgactgtgg 240  
 tctccaagac ccgcaagtac actgaagtga agccagccct cacaacatg gttgccgcag 300  
 caaaactcat c 311



<210> 2863  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2863  
  
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 aagtatggtg ctggaattgg ccttggtgtc tatgacatcc actccccaag aataccacca 120  
 actgaagaaa ttgctgacag aatcaacaag atgctggcag tgctcgagaa gaacatcttg 180  
 tgggtgaacc ctgactgtgg gctcaagacc cgtaagtaca ctgaggtgaa gccagccctc 240  
 acaaacatgg ttnnccgacg aaaactcatc cgcaacgaac ttgccaagtg anggta 296

<210> 2864  
 <211> 305  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2864  
  
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 ccagaggtaa tgctaccgtg cctgctatgg agatgancaa ntggttcgac accaactacc 120  
 actntattgt ccctgaattg ggccctgatg tgaacttcac ctatgcttct cacaaggctg 180  
 ttgatgaata caaggaggcc aaggcgcttg gagtggatac cattcccgtc ctcggtggcc 240  
 ctgttacata cttgttgctc tccaagcctg ccaagggagt cgagaaatcc ttttctctcc 300  
 tctct 305

<210> 2865  
 <211> 280  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2865  
  
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 actccatcat cgacatggac gctgatgtta tcaccattga gaactctcgc tccgatgaga 120  
 agtcctgtc agtcttccgt gaagggtgtga agtatggtgc tggaattggc cctggtgtct 180  
 atgacatcca ctccccaaga ataccaccaa ctgaagaaat cgctgacaga atcaataaga 240

tgcttgagct gctcgagaag aacatcttgt ggggtcaaccc 280

<210> 2866  
<211> 287  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2866

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ctggtaaatgc tgctgctctg gttcaagga agtcttctcc aagagtgacc aacgaggctg 120  
ttcagaaggc tgctgctgca ttgaagggtt cagatcatcg ccgtgcaaca atgtcagtg 180  
ccagactgga ttctcaacaa aagaagctca acctccaat cctgccaacc accactattg 240  
gacccctccc tcagactgta gaactgagga gggtaacgag tgaattc 287

<210> 2867  
<211> 324  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2867

ncntgcgtac gtaagctcgg nntncggnng ctcgngaatt cggctcgagc aagaagagct 60  
tgatattgat gttcttggtc atggagaacc agagagaaat gatatgggtg agtacttcgg 120  
tgaacaattg tncaggcttt gccttcaccg ttaatgggtg ggtgcaatcc tatgggtccc 180  
gttgcgtaga gccacgatca tctatggtga tgtgagccgc ccaaagccaa tgaccgtctt 240  
ctggatcatc ctggctcaga gctttaccaa ggcgccaatg aagggaatgc ttaccggtcc 300  
tgttaccatt ctcaactggc cctt 324

<210> 2868  
<211> 273  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2868

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atatgggtga gtacttcggt gagcaattgt caggctttgc cttcactggt aatgggtggg 120  
tgcaatccta tgggtcccggt tgtgtgaagc caccaatcat ctatgggtgat gtagccgccc 180

aaagccaatg actgtcttct ggtcatctct ggctcagagc tttaccaagc gcccaatgaa 240  
 gggaatgctt accggtcctg ttaccattct caa 273

<210> 2869  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2869

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 aagggttcag atcatcgccg tgcaacaaat gtcagtgcc aactggatgc tcaacaaaag 120  
 aagctcaacc ttccaatcct tccaaccacc actattggat ccttccctca gactgtagaa 180  
 ctgaggaggg tacgccgtga gttcaaggct aacaagatct ccgaggaaga gtatgttaag 240  
 tcaattaagg aggaaattcg caaagttggt gaacttcaag aagagcttga tattga 296

<210> 2870  
 <211> 301  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2870

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 agtcttctcc aagagtgacc aacgaggctg ttcagaaggc tgctgctgca ttgaagggtt 120  
 cagatcatcg ccgtgcaaca aatgtcagt ccagactgga ttctcaacaa aagaagctca 180  
 accttccaat cctgccaacc accactattg gatccttccc tcagactgta gaactgagga 240  
 gggtacgccg tgaattcaag gctaacaaga tctccgagga agagtatgta aagtcaatta 300  
 a 301

<210> 2871  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2871

cgtcgcangc acgcgtacgt nagctcgga ttcggctcga ngcagaactt gcgcctgctt 60  
 tgtctgggtt gaatgttctt gttgagacct actttgctga catccttgcg gaggcataca 120

agaccctcac atctctgaat ggcgtcactg catatggatt tnatttggtc cgtggaacca 180  
acactcttga tttgatcaag ggtggatttc ccagcgaaa atacctcttt gctggagtgg 240  
ttgatggaag gaacatctgg gccaatgacc ttgctgcttc tctcactacc ttgcagggtc 300

<210> 2872  
<211> 558  
<212> nucleic acid  
<213> Glycine max  
<400> 2872

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angcgtacgc nacgcntcc agagtgaacca acgangctgt tcataaagct gctgctgcat 120  
tgaangactg ggatggntc cgtgcaacaa atgtcagtgc cagactggat tctcaacana 180  
agaagctcaa ccttccaatc ctgccaacca ccactattgg atccttncct cagactgtat 240  
aactgaggag ggtacnccnt gaattnaagg ctaacangat ctccnaggaa nagtatgtaa 300  
agtcaattaa ngaggaaatt cgcaaanntt gtttnaactn naanaagagc ttgatattga 360  
tggtcttgtt catggatanc canagagaaa tgatatgggt gagtnctttn ggtgaacaaa 420  
ttttnaangc ttttnccctt taccgntaa tnggttggt gcaatnctat nggtttcccn 480  
ttgnngttaa agcctcctat cattttattg gngnttttta gccntccaa angccaattg 540  
accntcttt ttnttatt 558

<210> 2873  
<211> 279  
<212> nucleic acid  
<213> Glycine max  
<400> 2873

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ggtgagcaat tgtccaggct ttgccttcac tgtaaatggg tgggtgcaat cctatgggtc 120  
ccgttgtgtg aagccaccaa tcatctatgg tgatgtgagc cgcccaaagc caatgactgt 180  
cttctggtea tctctggctc agagctttac caagcgccca atgaaggga tgcttaccgg 240  
tcctgttacc attctcaatg gtcctttgtt agaaatgac 279

<210> 2874

<211> 295  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2874  
  
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 ctttgtctat caaggacgaa gtggaagacc ttgaaaaggc tggcatcact gttatccaaa 120  
 ttgatgaagc tgctttgaga gagggctctc cactgaggan atcagancaa gctcactact 180  
 tggactgggc tgtccatgcc ttcagaatca ccaatgttgg tgtccaggat accacccaga 240  
 tccacactca catgtgtctac tcgaattcaa cgacatcatc cactccatca tcgag 295

<210> 2875  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2875  
  
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 tgctgacctt aaggcagctg gtgcttcatt gattcagttt gatgagccta cccttgtctt 120  
 ggaccttgag tctcacaagt tgcaagcatt cactgacgca tatgcagaac ttgcgcctgc 180  
 tttgtctggg ttgaatgttc ttgttgagac ctactttgct gacatccctg ctgaggcata 240  
 caagaccctc acatctctga atggcgtcac tgcatatgga tttgatttgg tccgtggaac 300  
 caa 303

<210> 2876  
 <211> 293  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2876  
  
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 gtgctactcg aacttcaacg acatcatcca ctccatcatc gacatggacg ccgatgttat 120  
 caccattgag aactctcgct ccgacgagaa gctnctgtca gtcttccgcy aaggtgtgaa 180  
 gtatggtgct ggaattggcc ctggtgtcta tgacatccan tccccaaaga taccaccaac 240  
 tgangaaatt gctgacagaa tcaacaagat gctggcantg ctcgagaaga aca 293

<210> 2877  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2877  
  
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 tagatctgag accacctacc agattgcttt ggctatcaag gacgaagtgg aggaccttga 120  
 aaaggctggc atcactgtta tccaaattga tgaagctgct ttgagagagg gtctgccact 180  
 gaggaaatca gaacaagctc actacttgga ctgggctgtc catgccttca gaatcaccaa 240  
 tgttggtgtg caggatacca ctcagatcca caccacatg tgctactcca a 291

<210> 2878  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2878  
  
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 cggctcctgtt accattctca actggtcctt tgtagaaaat gaccaacctt gatctgagac 120  
 cacctaccag attgctttgt ctatcaagga cgaagtggaa gaccttgaaa aggctggcat 180  
 cactgttatc caaattgatg aagctgcttt gagagagggt cttccactga ggaaatcaga 240  
 gcaagctcac tacttggact gggctgtcca tgccctcaga atcaccaatg ttggtgtc 298

<210> 2879  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2879  
  
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 tggttcncgt tgtgtgaagc caccaatcat ctatggtgat gtgagcngnc caaagccaat 120  
 gactgtcttc tggctcatctc tggctcagag ctttaccaag cgcccaatga agggaatgct 180  
 taccggtcct gttaccattc tcaactggtc ctttggttaga aatgaccaac ctagatctga 240  
 gaccacctac cagattgctt tggctatcaa ggacgaagtg ggaggacctt gaaaaggctg 300

gcatcatggt

310

<210> 2880  
<211> 279  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2880

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gatgtgagcc gcccaaagcc aatgactgtc ttctgggtcat ctctgggtca gagctttacc 120  
aagcgcccaa tgaagggaat gcttaccggt cctgttacca ttctcaactg gtcccttntt 180  
agaaatgacc aacctagatc tgagaccacc taccagattg ctttggtat caaggacgaa 240  
gtggaggacc ttgaaaaggc tggcatcact gttatccaa 279

<210> 2881  
<211> 280  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2881

tttatgcagc gtacgtnagc tcggaattcg gctcgaggnc aattaaggag gaaattcgca 60  
aagttgttga acttcaagaa gagcttgata ttgatgttct tgttcatgga gaaccagaga 120  
gaaatgatat ggttgagtac ttcggtgagc aattgtcagg cttgccttca ctgttaatgg 180  
gtgggtgcaa tcctatggtt cccgttgtgt gaagccacca atcatctatg gtgatgtgag 240  
ccgccccaaag ccaatgactg tcttctgggtc atctctggct 280

<210> 2882  
<211> 344  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2882

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ctcagactgt agaactgagg agggtagcc gtgaattcaa ggctaacaag atctccgagg 120  
aagagtatgt aaagtcaatt aaggaggaaa ttcgcaaagt tggtgagctt caagaagagc 180  
ttgatattga tggtcttggt catggagaac cagagagaaa tgatatggtt gagtacttcg 240

gtgaacaatt gtcaggcttg ccttcaccgt taatgggtgg gtgcaatcct aggttcccgt 300  
 tgcgtgaagc caccgatcat ctatggtgag tgagccgccc aaag 344

<210> 2883  
 <211> 276  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2883

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 gagctttacc aagcgcccaa tgaagggaat gcttaccggt cctggtacca ttctcaactg 180  
 gtcctttggt agaaatgacc aacctagatc tgagaccacc taccagattg ctttgtctat 240  
 caaggacgaa gtggaagacc ttgaaaaggc tggcat 276

<210> 2884  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2884

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 actgtcttct ggtcatctct ggctcagagc ttaccaagc gcccaatgaa gggaatgctt 180  
 accggtcctg ttaccattct caactggtcc tttgttagaa atgaccaacc tagatctgag 240  
 accacctacc agattgcttt ggctatcaag gacgaatgga ggaccttg 288

<210> 2885  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2885

nngcangcac gcntacgtna gctcgggaatt cggctcgagn gaccttgctg cttctctcac 60  
 tacattgnca gggctctgag ggcatgtgg gcaaagataa gcttggtgtg tccacctcct 120  
 cctcccttct tcacactgct gttgatcttg ttaacgagac caagttggat gacgagatca 180



agtcatggct agcatttgct gcacaaaaaa ttgttgaagt taacgcattg gctaaggcat 240  
 tgtctggcaa caaggatgtg gccttcttct ctgctaatac tgcagctcag gcttcaagga 300  
 agtcctctcc aaga 314

<210> 2886  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2886

gnogcatgca cgcgtacgta agctcggaat tcggctcgag cttgatttga tcaaggggtgg 60  
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 tgaccttgcg gcttctctca ctaccttgca gggctctgag ggcattgtgg gcaaagataa 180  
 gcttggttgg tccacctcct cctcccttct tcacactgct gttgatctag ttaacgagac 240  
 caagttggat gatgagatca agtcatggct agcttttgcg gcccaaaaaa tgttgaagtt 300  
 aacg 304

<210> 2887  
 <211> 275  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2887

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 ttctgggtcat ctctggctca gagctttacc aagcgcccaa tgaagggaat gcttaccggt 120  
 cctgttacca ttctcaactg gtcctttggt agaaatgacc aacctagatc tgagaccacc 180  
 taccagattg ctttggctat caaggacgaa gtggaggacc ttgaaaaggc tggcatcact 240  
 gttatccaaa ttgatgaagc tgctttgaga gaggg 275

<210> 2888  
 <211> 257  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2888

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tgctgacatc cctgctgagg cgtacaagac cctcacatct ctgaatggcg tcaactgcata 120  
 tgggtttgat ttggtccgtg gaaccatac tcttgatttg atcaaggggtg gatttcccag 180  
 tggaaaatac ctctttgctg gagtgggtga tggaaggaac atctgggcca atgaccttgc 240  
 tgcttctctc actacat 257

<210> 2889  
 <211> 278  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2889

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 tggatttgat ttggtccgtg gaaccaaacac tcttgatttg atcaaggggtg gatttcccag 180  
 cggaaaatac ctctttgctg gagtgggtga tggaaggaac atctgggcca atgaccttgc 240  
 tgcttctctc actaccttgc agggctcttga gggcattg 278

<210> 2890  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2890

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 gaggaaatca gagcaagctc actacttggc ctgggctgtc catgccttca gaatcaccaa 180  
 tgttggtgtc caggatacca cccagatcca cactcacatg tgctactcga acttcaacga 240  
 catcatccac tccatcatcg acatggacgc cgatgttata accatgagaa tctcgtc 298

<210> 2891  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2891

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ntccgaggaa gagtatgtta 60

agtcaattaa ggaggaaatt cgcaaagttg ttagaacttc aagaagagct tgatattgat 120  
 gttcttggtc atggagaacc agagagaaat gatatggttg agtacttcgg tgagcaattg 180  
 tnangctttg ctttactgt taatgggtgg gtgcaatcct atggttcccg ttgtgtgaag 240  
 ccaccaatca tctatggtga tgtgagccgc ccaaagccaa tgactgtctt ctggtcattt 300  
 ctggctcaga gcttta 316

<210> 2892  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2892

gtgcgancga cgcntacgtn agctcgggaa ttcggctcga gngactggat gctcaacaaa 60  
 agaagctcaa cttccaatc cttccaacca nactattgg atccttccct cagactgtag 120  
 aactgaggag ggtacgccgt gagttcaagg ctaacaagat ctccgaggaa gagtatgtta 180  
 agtcaattaa ggaggaaatt cgcaaagttg ttgaacttca agaagagctt gatattgatg 240  
 ttcttggttca ggagaaccag agagaaatga tatgggttgag tacttcggt 289

<210> 2893  
 <211> 320  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2893

gttactgcac gngtacgtaa gctcgggaatt nggctcgagc tcgagccgct atggtgatgt 60  
 gagccgcca aagccaatga ccgtcttctg gtcattcttg gtcagagct ttaccaagcg 120  
 cccaatgaag ggaatgctta ccggtcctgt taccattctc aactggctct ttgttagaaa 180  
 tgaccaacct agatctgaga ccacntacca gattgctttg tctatcaagg acgaagtgga 240  
 agaccttgaa aaggctggca tctactgttat ccaaattgat gaagcgcttt gagagagggt 300  
 cntccactga ggaaatcaga 320

<210> 2894  
 <211> 302  
 <212> nucleic acid  
 <213> Glycine max

<400> 2894

gtcgcacatgca cgcgtacgta agctcgggaat tcggctcgag cgtaaatggg tgggtgcaat 60  
cctatgggttc ccgttgctgtg aagccaccga tcctctatgg tgatgtgagc cgcccaaagc 120  
caatgaccgt cttctgggtca tctctggctc agagntttac caagcgccca atgaagggaa 180  
tgcttaccgg tcctgttacc attctcaact ggtcctttgt tagaaatgac caacctagat 240  
ctgagaccac ctaccagatt gctttgtcta tcaaggacga atggaagacc ttgaaaaggc 300  
tg 302

<210> 2895

<211> 313

<212> nucleic acid

<213> Glycine max

<400> 2895

acgtcgcacatg cagcgtacg tnagctcgga attcggctcg agnttaatgg gtgggtgcaa 60  
tcctatgggtt ccggttgctg gaagccaccg atcatctatg gtgatgtgag ccgccccaaag 120  
ccaatgaccg tcttctggctc atctctggct cagagcttta ccaagcgccc aatgaaggga 180  
ntgcttaccg gtctgtttac cattctcaac tggctcctttg ttagaaatga ccaacctaga 240  
tctgagacca cctaccagat tgctttgtct atcaaggacg aatggaagac cttgaaaagg 300  
ctggcatcat gtt 313

<210> 2896

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2896

cantcgnang nacgcgtacg taagctcgga attcgggtnc naggaaanta cctctttgct 60  
ggagtgggtg atggaaggaa catctgggcc aatgaccttg ctgcttctct cactaccttg 120  
cagggtcttg agggcattgt gggcaaagat aagcttggtg tgtccacctc ctctccctt 180  
cttcacactg ctgttgatct agttaacgag accaagttgg atgatgagat caagtcatgg 240  
ctagcttttg ctgccccaaa aattgttgaa gttaacgcac ggctaaagca tgtctggcca 300  
caaggatgag gg 312

<210> 2897  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2897  
  
 gcngcacgcg tacgtaagct cggaattcgg ctcgaggatt gcacccaccc attaacggtg 60  
 atgtgagccg cccaaagcca atgaccgtct tctggtcac tctggctcag agctttacca 120  
 agcgcccaat gaagggaatg cttaccggtc ctgttaccat tctcaactgg tcctttgtta 180  
 gaaatgacca acctagatct gagaccacct accagattgc tttgtctatc aaggacgaag 240  
 tggaagacct tgaaaaggct ggcatactg ttatccaaat tgatgaagct g 291

<210> 2898  
 <211> 312  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2898  
  
 agtcgcatgc acgcgtacgt aagctcggaa ttcggtcga gtcagatgt tenttccatc 60  
 aaccactcca gcnaagataa ncttgtngtg tccacctcct cctcnnttct tcacactgct 120  
 gtngatnttg ttaacgagac caagttggat gacgagatca agtcattggt agcatttgct 180  
 gcacaaaaaa ttgttgaagt taacgnattg gctaaggcnt tgtctggcaa caaggatgtg 240  
 gccttcttnt ctgctaattgc tgcagctcag gcttcaagga agtcntctcc aagagtgacc 300  
 aacgaggctg tt 312

<210> 2899  
 <211> 247  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2899  
  
 cgcatatgca gaacttgac ctgctttgtc tgatctgaat gtncctgttg anacctactt 60  
 tgctgacatc cctgctgagg cgtacaagac cctcacatct ctgaatggcg tcaactgcata 120  
 tgggtttgat ttgggtccgtg gaaccatac tottgatttg atcaagggtg gatttcccag 180  
 tggaaaatac ctctttgctg gagtggttga tggaaggaac atctgggcca atgaccttgc 240

tgcttct 247

<210> 2900  
<211> 317  
<212> nucleic acid  
<213> Glycine max

<400> 2900

tgcacncaac gcgtacgtaa gctcggaatt cggctcgagg ggaatgctta ccggctcctgt 60  
taccattctc aactggctct ttgttagaaa tgaccaacct agatctgaga ccacctacca 120  
gattgctttg tctatcaagg acganacngg aagaccttga aaaggctggc anccactggt 180  
ntccaaattg atgaagctgc tttagagagag ggtcttccac tgaggaaatc agagcaagct 240  
cactacttgg actgggctgt ccatgccttc agaataacca atgttggtgt ccaggataacc 300  
accagatcc acactna 317

<210> 2901  
<211> 285  
<212> nucleic acid  
<213> Glycine max

<400> 2901

gtctatgcac gcgtacgtaa gctcggaatt cggctcgagc gagaccangt tggatgacga 60  
gatcaagtca tggctagcat ttgctgcaca aaaaattggt gaagttaacg cattggctaa 120  
ggcattgtct gncaacaagg atgtggcttc ttctctgcta atgctgcagc tcaggcttca 180  
aggaagtcct ctccaagagt gaccaacgag gctgttcaga aggctgctgc tgcattgaag 240  
ggttcagatc atcgccgtgc aacaaatgtc agtgccagac tggat 285

<210> 2902  
<211> 264  
<212> nucleic acid  
<213> Glycine max

<400> 2902

gtcgacngca cgcgtacgta agctcggaat tcggctcgag nacaagatct ccgaggaaga 60  
gtatgtaaag tcaattaagg aggaaattcg caaagttggt gagcttcaag aagagcttga 120  
tattgatgtt cttgttcatt gagaaccaga gagaaatgat atggttgagt acttcggtga 180

acaattgtca ggctttgect tcaccgttaa tgggtgggtg caatcctatg gttcccgttg 240  
cgtgaagcca ccgatcatct atgg 264

<210> 2903  
<211> 299  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2903

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctgaattggg ccctgatgtg 60  
aacttcacct atgcttctca caaggctgtt gatgaatata aggaggccaa ggcgcttgga 120  
gtggatacca ttcccgtact cgttggccct gttacatact tgttgctctc caagcctgcc 180  
aagggagtcg agaaatcctt ttctctctc tctctccttc ccaaggttct tgctgtctac 240  
aaggaagtta ttgctgacct taaggcagct ggtgcttcat ggattcaatt gatgagcct 299

<210> 2904  
<211> 305  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2904

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ggcagctggg gcttcattgn ttcaatttga tgagcctacc cttgtcttgg accttgaatc 120  
tcacaagttg caagctttca ctgacgcata tgcanaactt gcacctgctt tgtctgatct 180  
gaatgttctt gttgagacct actttgctga catccctgct gaggcgtaca agaccctcac 240  
atctctgaat ggcgtcactg catatgggtt tgatttggtc cgtggaaccc atactcttga 300  
tttga 305

<210> 2905  
<211> 299  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2905

actacgcatg cacgcgtacg taagctcgga attcggctcg agtctggctc agagctttac 60  
caagcgccca atgaaggga tgcttacncc tctgtttacc attctncaac tggtcctttg 120

ttagaaatga ccaacctaga tctgagacca cctaccagat tgctttgtct atcaagntga 180  
 agtggaagac cttgaaaagg ctggcatcac tgttatccaa attgatgaag ctgctttgag 240  
 agaggggtctt ccaactgagga aatcagagca agctcactac ttggactggg ctgtccatg 299

<210> 2906  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2906

cacgcangca cgcgtagcta agctcggaat tcggctcgag gtggatttcc cagtggaaaa 60  
 tacctctttg ctggagtggg tgatggaagg aacatctggg ccaatgacct tgctgcttct 120  
 ctcactacat tgcaggggtct tgagggcatt gtgggcaaag ataagcttgt tgtgtccacc 180  
 tcccctccct tcttcacact gctgttgatc ttgttaacga gaccaagttg gatgacgaga 240  
 tcaagtcatg gctagcattt gctgcacaaa aaattgttga agttaa 286

<210> 2907  
 <211> 313  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2907

agtcgctgcn nncgtcgga ttcggctcga ggtctacaag gaagtnattg ctggacctta 60  
 aggcagctgg tgcttcatgg attcaatntg atgagcctac ccttatcttg gaccttgaat 120  
 ctcanaagtt gcaagctttc actgacgcat atgcagaact tgcacctgct ttgtctgac 180  
 tgaatgtntc ngtnagacn cactttgctg acatccctgc tgaggcgtac aagaccctca 240  
 catctctgaa tggcgtcact gcatatgggt ttgatttggt ccgtggaacc catactcttg 300  
 atttgatcaa ggg 313

<210> 2908  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2908

acgtcgcan cacgcgtacg taagctcgga attcggctcg agtggattct caacaaaaga 60



agctcaacct tccaatcctt ccaaccacca ctattggntc cttccctcag actgtagaac 120  
 tgaggagggt acgccgtgaa ttcaaggcta acaagatctc cgaggaagag tatgtaaagt 180  
 caattaagga ggaaattcgc aaagttgttg agcttcaaga agagcttgat attgatgttc 240  
 ttgttcatgg agaaccagag agaaatgata tgggt 274

<210> 2909  
 <211> 276  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2909

gcatgcncgc gtncgtnagc tcggaattcg gctcgagtct ggctcagagc tttaccaagc 60  
 gcccaatgaa gggaatgctt accggctctg ttaccattct caactggctc tttgttagaa 120  
 atgaccaacc tagatctgag accacctacc agattgcttt gtctatcaag gaogaagtgg 180  
 aagaccttga aaaggctggc atcactgtta tccaaattga tgaagctgct ttgagagagg 240  
 gtcttccact gaggaaatca gagcaagctc atactt 276

<210> 2910  
 <211> 252  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2910

cacaagtngc aagcnttcac tgacgcatat gcagaacttg cacctgcttt gtctgatctg 60  
 aatgttcttg ttgagacctt ctttgctgac atccctgctg aggcgtacaa gaccctcaca 120  
 tctctgaatg gcgtcactgc atatgggttt gatttggctc gtggaaccca tactcttgat 180  
 ttgatcaagg gtggatttcc cagtggaaaa tacctctttg ctggagtggg tgatggnagg 240  
 nacatctggg cc 252

<210> 2911  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2911

gtcgcangca cgcgtacgta agctcggaat tcggctcgag attggctaag gcattgtctg 60

gcaacaagga tgtggccttc ttctctgcta atgctgcagc ttcaggcttc aaggaagtcc 120  
tctccaagag tgaccaacga ggctgttcag aaggctgctg ctgcattgaa gggttcagat 180  
catcgccgtg caacaaatgt cagtgccaga ctggntgctc aacaaaagaa gctcaacctt 240  
ccaatccttc caaccaccac tattgntcct tcctcagacg tgtgaa 286

<210> 2912  
<211> 293  
<212> nucleic acid  
<213> Glycine max  
<400> 2912

cgtcgnngca cgcgtacgtn agctcggnan tnggctcgag ctcgagccgc aagagtgacc 60  
aacgaggctg ttcagaaggc tgctgctgca ttgaaggggt cngatcatcg ccgtgcaaca 120  
aatgtcagtg ccagactgga ttctcaacan aagaagctca accttccaat cctgccaaac 180  
accactattg gatccttccc tcagactgta gaactgagga gggtagcccg tgaattcnag 240  
gctaacaaga tctccgagga agagtatgta nngtcaatta agngngaaat tcg 293

<210> 2913  
<211> 274  
<212> nucleic acid  
<213> Glycine max  
<400> 2913

gtcnaatgca cgcgtacgta agctcggaat tcggctcgag cgaggctggt cagaaggctg 60  
ctgctgcatt gaaggggttca gatcatcgcc gtgcaacaaa tgtcagtgcc agactggatt 120  
ctcaacaaaa gaagctcaac ctccaatcc tgccaaccac cactattgga tccttccttc 180  
agactgtaga actgaggagg gtacgccgtg aattcaaggc taacaagatc tccgaggaag 240  
agtatgtaaa gtcaattaag gaggaaattc gcaa 274

<210> 2914  
<211> 283  
<212> nucleic acid  
<213> Glycine max  
<400> 2914

acantcatgc acgcgnagta gtcggaatnc ggctcgaggt tgagcnncaa gnagancttg 60

atattgnngt tcntgntcat ggaganccan agagaaatga tatggttgag tacttcgggtg 120  
aacaattgtc aggcttttgct ttcaccgtta atgggtgggt gcaatcctat ggntcccgtt 180  
gcgtgaagcc ancgatcatc tatggtgatg tnagccgccc aaagccaatg accgtntttct 240  
ggtcattctct ggctcagagc tttaccaagc gccaatgaag gga 283

<210> 2915  
<211> 534  
<212> nucleic acid  
<213> Glycine max  
<400> 2915

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gagggattnn aaantctttc nanagccatg ttgtncocgt ttcggaattc ccgctcgacc 60  
cacgcgtccg nncacnctc cgctgcagaa gacgactgaa nggagagctt gatattgatg 120  
ttcttgggca tggagaacca gagagaantg atgtggttga gtacttcgggt gagcaattgt 180  
caggctttgc cttcactgtt aatgggtggg tgcaatccta tggttcccgt tgtgtgaanc 240  
caccnatcat ctatggtgat gtgagccccc aaanccaatg actgtcttct ggtcatctct 300  
ggctcananc tttaccaatc gcccnntgaa anggaatnct tnccgggocct gttacattnt 360  
naacttgggc cttnttttna anatancaa cctatttntt annccnctnc nttattnttt 420  
tnncttttna ggatnatnng nttgnntttt tanaaanggg ttngatnat tnttttntn 480  
natnttnttn atnnnnntn tnntnaaang ntntntnat tnngnaataa natt 534

<210> 2916  
<211> 297  
<212> nucleic acid  
<213> Glycine max  
<400> 2916

gttaacgaga ccaagcnanc catgacttga tctcatcatc caacttggtc tcgttaacta 60  
gatcaacagc agtgtgaaga agggaggagg agatcaagtc atggctagct tttgctgccc 120  
aaaaaattgt tgaagttaac gcattggcta aagcattgtc tggccacaag gatgaggcct 180  
tcttctctng taatgtctgt nctctggctt caaggaagtc ttctccaaga gtgaccaacg 240  
aggctgttca gaaggctgct gctgcattga agggttcaga tcatcgccgt gcaacaa 297

<210> 2917

<211> 410  
 <212> nucleic acid  
 <213> Glycine max  
  
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 gtatggtgct ggaattggcc ctggtgtcta tgacatccac tccccaaaga taccaccaac 60  
 tgaagaaatc gctgacagaa tcaataagat gcttgcagtg ctogagaaga acatcttgtg 120  
 ggtcaaccct gactgtggtc ttaagacccg caagtacact gaagtgaagc cagccctcac 180  
 aaacatgggt gccgcagcaa aactcatccg taacgaactt gccaaagtga tgggtataaga 240  
 aagtagaatc tacaagttca ttggttctgc ttttataata caccaaagaa aaattttcta 300  
 tattgggttg tttcaataac cgtgtgtgga atatttagat gttttagcat gctccgtgaa 360  
 caattgatcc tcctcaaacc ctctcccctt aattttccca actcccgtt 410

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ggctggcatc actgttatcc agnttgggtga agctgctttg agagaggggtc ttccactgag 240  
gaaatcagan caagctcact attggatggg tgtccatgcc ttcagaatca ccangttggg 300  
gtccaggata cca 313

<210> 2920  
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<213> Glycine max  
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attggctaag gcattgtctg gcaacaagga tgtggccttc ttctctgcta atgctgcagc 120  
tcaggcttca aggaagtctc ctccaagagt gaccaacgag gctgttcaga aggctgctgc 180  
tgcattnaag ggttcagntc ancgncgtnc aacanntcnc agccnnantg ganantcncn 240  
aaaaaaggct cncncncc 259

<210> 2921  
<211> 286  
<212> nucleic acid  
<213> Glycine max  
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tcgctgcacg cgtacgttag ctcggaattc ggctcgaggg ataccgttcc ggtcctcggt 60  
ggccctgtta catacctgtt gctctccaag cctgccaaagg gagttgagaa atccttttct 120  
ctcctctctc tccttcccaa ggttcttgct gtctacaagg aagttattgc tgaccttaag 180  
gcagctggtg cttcatggat tcagtttgat gagcctaccc ttgtcttgga ccttgagtct 240  
cacaagttgc aagcattcac tgacgcatat gcagaacttg cgctg 286

<210> 2922  
<211> 242  
<212> nucleic acid  
<213> Glycine max  
<400> 2922

gtgcaatcct atgggtcccg ttgctgaag ccaccgatca tctatgggtga tgtgagccgc 60  
ccaaagccaa tgaccgtctt ctggctcatct ctggctcaga gctttaccaa gccccaatga 120

aggaatgct taccggctcct gttaccattc tcaactggtc ctttgtaga aatgaccaac 180  
ctagatctga gaccacctac cagattgctt tgtctatcaa ggacgaatgg aagacctga 240  
aa 242

<210> 2923  
<211> 270  
<212> nucleic acid  
<213> Glycine max  
<400> 2923

gogtacgtaa gctcggaatt cggctcgagt cagatcatcg ccgtgcaaca aatgtcagtg 60  
ccagactgga ttctcaacaa aagaagctca accttccaat cctgcccaacc accactattg 120  
gatacttccc tcagactgta gaactgagga gggtagccg tgaattcaag gctaacaaga 180  
tctccgagga agagtatgta agtcaattaa ggaggaaatt cgcaaagttg ttgagcttca 240  
agaagagctt gatattgatg ttcttgttca 270

<210> 2924  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
<400> 2924

agtcnnangc acgntacgt aagctcggaa ttccggctcga gggtaggattt ccagtgga 60  
aatacctctt tgctggagtg gttnatggaa ggaacatctn gngccaatga ccttgctgct 120  
tctctcacta cattgcaggg tcttgagggc attgtnggca aagataagct tgttgtgtcc 180  
acctctctct ccttcttcca cactgctgtt gatcttgta acgagaccaa gttggatgac 240  
gagatcaagt catggctagc atttgctgca caaaaaattg tgaagttaac gc 292

<210> 2925  
<211> 312  
<212> nucleic acid  
<213> Glycine max  
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aanatttnnc gtcgcatgca tgcgtacgta agctcggaat tcggctcgag gagaaatgat 60  
atggttctcc atgaacaaga acatcaatat caagctcttc ttgaagttct tgttcatgga 120

gaaccagaga gacnatgata tgggtgagta cttcgggtgag caattgtcag gctttgcctt 180  
 cactgttaat ggggtgggtnc natectatgg ttcccgttgt gtgaagccac caatcatcta 240  
 tggatgatgtg agcgcgcccac agccaatgac tgtcttcttg tcctctcttg ctcagagctt 300  
 taccaagcgc cc 312

<210> 2926  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max  
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tcgcangcac gcgtacgtna gctcgggaatt cggctcgagc aagtgggttcg acaccaacta 60  
 ccaactgggat tgcctctgaa ttgggcccctg atgtgaactt cacctatgct tctcacaagg 120  
 ctgttgatga atacaaggag gccaaaggcgc ttggagtggg taccattccc gtactcggtt 180  
 gccctgttac atacttggtg ctctccaagc ctgccaaggg agtcgagaaa tctttttctc 240  
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<210> 2927  
 <211> 258  
 <212> nucleic acid  
 <213> Glycine max  
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 attctcaact ggtctttggt agaaaatgac caacctagat ctgagacaac taccagattg 180  
 ctttgtctat ccaaggacga ntnggaagac cttgaaaagg ctggcatcac tgttatccaa 240  
 attgatgaag ctgctttg 258

<210> 2928  
 <211> 335  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2928

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ctatgacatc cactcccca gaataccacc aactgaagaa atcgctgaca gaatcaataa 120  
gatgcttgca gtgctcgaga agaacatctt gtgggtcaac cctgactgtg gtctcaagac 180  
ccgcaagtac actgaagtga agccagccct cacaacatg gttgccgcag caaaactcat 240  
ccgtaacgaa cttgccaaagt gaatggtata agaaagtaga atctacaagt tcattgggtc 300  
tgcttttata atacaccaa gaaaaatttt ctata 335

<210> 2929  
<211> 279  
<212> nucleic acid  
<213> Glycine max  
<400> 2929

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annntangta cgcgtacgta agctcggaat tccgctcgag caccaactac cactttattg 60  
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acaaggnggc caaggcgctt ggagtggata ccattcccgt actcgttggc cctgttacat 180  
acttgttgct ctccaagcct gccaaaggag tcgagaaaac cttttctctc ctctctctcc 240  
ttcccaaggt tcttgctgtc tacaaggaag ttattgctg 279

<210> 2930  
<211> 282  
<212> nucleic acid  
<213> Glycine max  
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gtcgcangca cgcgtacgtn agctcggaat tccgctcgag nctcgctcc gatgagaagc 60  
tcctgtcagt cttccgtgaa ggtgtgaagt atgggtgctg aattggccct ggtgtctatg 120  
acatccactc cccaagaata ccaccaactg aagaaatcgc tgacagaatc aatacgatgc 180  
ttgcagtgct cgagaagaac atcttggtgg tcaaccctga ctgtggtctc aagaccgcga 240  
agtacactga atgaagccag cctcacaaa catggttgcc gg 282

<210> 2931  
<211> 261  
<212> nucleic acid  
<213> Glycine max  
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ccttccaatc cttccaacca ccactattgg ntccttccct cagactgtag aactgaggag 60  
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 aggaggaaat tcgcaaagtt gttgaacttc aagaagagct tgatattgat gttcttgttc 180  
 atggagaacc agagagaaat gatatggttg agtacttcgg tgagcattgt caggctttgc 240  
 ctcaactgtta atgggtgggt g 261

<210> 2932  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
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ctcccnacg tngcatgcac gcgtacgtna gctcngaat tcggctcgag gaaatccttt 60  
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 aaggcagctg gtgcttcatg gattcaattt gatgagccta cccttgtctt ggaccttgaa 180  
 tctcacaagt tgcaagcttt cactgacgca tatgcagaac ttgcacctgc tttgtctgat 240  
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<210> 2933  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max  
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tcgcangcac gcgtacgtna gctcggaatt cggtcgagc ttgttgctct ccaagcctgc 60  
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 caaggaagtt attgctgacc ttaaggcagc tgggtgcttca tggattcaat ttgatgagcc 180  
 tacccttgtc ttggaccttg aatctcaciaa gttgcaagct ttcactgacg catatgcaga 240  
 acttgcacct gctttgtctg atctgaatgt tcttgttgag actactttgc tgacatcc 298

<210> 2934  
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 <212> nucleic acid  
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agtcgcangc acgcgtacgt aagctcggaa ttcggctcga gcaaggcgct tggagtggat 60  
accattcccc tactcgttgg ccctgttaca tacttggtgc totccaagcc tgccaaggga 120  
gtcgagaaat ctttttctct cctctctctc cttcccaagg ttotttctgt ctacaaggaa 180  
gttattgctg accttaaggc agctggtgct tcatggattc aatttgatga gcctaccctt 240  
gtcttggacc ttgaatctca caagttgcn 269

<210> 2935  
<211> 261  
<212> nucleic acid  
<213> Glycine max  
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aattgctgac agaatcaaca agatgctggc agtgctcgag aagaacatct tgtgggtgaa 120  
ccctgactgt gggctcaaga cccgtaagta cactgaggtg aagccagccc tcacaaacat 180  
ggttgccgca gcaaaactca tccgcaacga acttgccaag tgaatggtat aagaaagtag 240  
aatcttccaa gtcatttgggt t 261

<210> 2936  
<211> 262  
<212> nucleic acid  
<213> Glycine max  
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agcgcccaat gaagggaatg cttaccggtc ctgttaccat tcncaactgg tcctttgtta 120  
gaaatgacca acctagatct gagaccacct accagattgc tttgtctatc aaggacgaag 180  
tggaagacct tgaaaaggct ggcactcactg ttatocaaat tgatgaagct gctttgagag 240  
agggtctcca ctgaggaaat ca 262

<210> 2937  
<211> 280  
<212> nucleic acid  
<213> Glycine max  
<400> 2937

acgcgtacgt aagctcggaa ttcggctcga ggtgaagtat ggtgctggaa ttggccctgg 60  
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 caagatgctg gcagtgctcg agaagaacat cttgtgggtg aaccctgact gtggggtcaa 180  
 gacccgtaag tacactgagg tgaagccagc cctcacaaac atggttgccg cagcaaaaact 240  
 ncatccgcaa cgaattgcc aatgatggtat aagaaaataga 280

<210> 2938  
 <211> 244  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2938

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 gttgatggaa ggaacatctg ggccaatgac cttgctgctt ctctcactac attgcagggt 180  
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 ctgc 244

<210> 2939  
 <211> 289  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2939

tctatgcacg cgtacgnag ctcggaattc ggctcgagca atgaactgtct tctgggtcatc 60  
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 tctcaactgg tcctttgtta gaaatgacca acctagatct gagaccacct accagattgc 180  
 tttggctatc aaggacgaag tggaggacct gaaaaggctg gcatcactgt tatccaaatt 240  
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<210> 2940  
 <211> 301  
 <212> nucleic acid  
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 <400> 2940

ncgtcgcang cacgcgtacg tnagctcgga attcggctcg agggttcccg ttgcgtgaag 60  
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ctggctcaga gctttaccaa ggcgccaatg aagggaatgc ttaccggtcc tgttancatt 180  
ctcaactggt cctttgttag aaatgaccaa cctagatctn cagaccacct accagattgc 240  
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g 301

<210> 2941  
<211> 295  
<212> nucleic acid  
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<400> 2941

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tcgcatgcac gcgtacgtaa gctcggaatt cggctcgagc cagattgctt tggctatcaa 60  
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ccatgccttc agaatcacca atgttggtgt gcaggatacc actcagatcc acaccacat 240  
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<210> 2942  
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<212> nucleic acid  
<213> Glycine max

<400> 2942

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ttgatcaagg gtggatttcc cagtggaaaa tacctctttg ctggantggt tnatggangg 180  
nncatctggg ccaatgacct tgngcntct ctcachacat tncagggtct tgagggcatt 240  
gtgggcnaag atannctngt tgtgtccacc tctccnccc ttentcacac tgctg 295

<210> 2943  
<211> 269  
<212> nucleic acid  
<213> Glycine max

<400> 2943

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gctgggtgctt catggattca atttgatgag cctacccttg tcttggacct tgaatctcac 120  
aagttgcaag ctttcaactga cgcataatgca gaacttgacac ctgctttgtc tgatctgaat 180  
gttcttgttg agacctactt tgctgacatc cctgctgagg cgtacaagac cctcacatct 240  
ctgaatggcg tcaactgcata tgggtttga 269

<210> 2944

<211> 312

<212> nucleic acid

<213> Glycine max

<400> 2944

tgcgancac gcgtacgtaa gctcggaatt cggctcgagc tgggtgtctat gacatccact 60  
ccccagaat accaccaact gaagaaattg ctgacagaat caacaagatg ctggcagtg 120  
tcgagaagaa catcttgttg gtgaaccctg actgtgggct caagaccctg aagtacactg 180  
aggtgaagoc agcctcaca aacatggttg ccgcagcaaa actcatccgc aacgaacttg 240  
ccaagtgaat ggtataagaa agtagaatct tccaagtcac ttggttctgc tttatattat 300  
aatacaccaa ag 312

<210> 2945

<211> 320

<212> nucleic acid

<213> Glycine max

<400> 2945

gangcacgcg tacgtnagct cggaattcgg ctcgagccca gcggaaaata cctctttgct 60  
gncagtggtt gatggaagga acatctgggg caatggacct ttgtggnctt tctcactac 120  
cttggcaggg tcttganggg cattgtgggc aaagataagc ttgttggtgc cactcctcc 180  
tcccttcttc aactgtctgt tgaccagtt aacgagacca agttggatga tgagatcaag 240  
tcatggctag cttttgctgc ccaaaaaaat tgttgaagtt aacgcattgg ctaaagcatt 300  
gtctggccac aaggatgagg 320

<210> 2946

<211> 299  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2946  
  
 nnangcatgc acgcgtacgt aagctnngaa ttcggctcga gctgaatggn gtcactgnca 60  
 tatgggnttg atttgggtccg tggaacccat actgcttgat ttgatcaagg gtggatttcc 120  
 cagtggaaaa tacctctttg ctggagtggg tgatggaagg aacatctggg ccaatgacct 180  
 tgctgcttct ctactacat tgcagggctc tgagggcatt gtgggcaaag ataagcttgt 240  
 tgtgtccacc tcctctctcc ttcttcacac tgctgttgat ccttggttaac gagaccaag 299

<210> 2947  
 <211> 269  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2947  
  
 aataatatgc acgcgtacgt aagntcggaa ttcggctcga gacaagttgc aagcattcac 60  
 tgacgcntat gcagaacttg gcctgcttg ggggtggttn aatgtnttg ttgagaccga 120  
 ctttnttgac atccctgctg aggcatacaa gaccctcaca tctctgaatg gcgtcactgc 180  
 atatggattt gatttgggtc gtggaaccaa cactcttgat ttgatcaagg gtggatttcc 240  
 cancgaaaa tacctctttg ctggagtgg 269

<210> 2948  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2948  
  
 tcgcatgcac gcgtacgtaa gctcgggaat tcggctcgag attggatcct tccctcagac 60  
 tgtagaactg aggagggtac gccgtgagtt caaggctaac aagatctccg aggaagagta 120  
 tgttaagtca attaaggagg aaattcgcaa agttgttgaa cttcaagaag agctgatatt 180  
 gatgttctgt tcatggagaa ccagagagaa atgatatggg tgagtacttc ggtgagcaat 240  
 tgtcaggctt tgcttcactg ttaatgggtg ggtgcatact atggttcccg ttgt 294

<210> 2949

<211> 280  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2949  
 nacntngcat gcacgcgtac gtaagctcgg aattcggctc gaggttacca ttctcaactg 60  
 gtcctttgtt agaaatgacc aacctagatc tgagaccacc taccagattg ctttgtctnc 120  
 caaggacgaa gtggaagacc ttgaaaagnc tggcatcact gttatccaaa ttinatgaagc 180  
 tgctttgann gagggctctc cactnnggaa atcagagcan ntcactactt gganegggt 240  
 gtccatntnt tcagaatcac nnntgttggt gtccnngata 280

<210> 2950  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2950  
 atgencgcgt acgt nagctc ggaattcggc tcgagcattc ccgtactcgt tggccctggt 60  
 acatacttgt tgctctccaa gcttgccaag ggagtcgaga aatccttttc tctcctctct 120  
 ctctttccca aggttcttgc tgtctacaag gaagttattg ctgaccttaa ggcagctggt 180  
 gcttcatgga ttcaatttga tgagcctacc cttgtcttgg acttgaatct cacaagttgc 240  
 aagctttcat gacgcatatg cagaattgca ctgt 274

<210> 2951  
 <211> 270  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2951  
 ncgtcgcattg cacgcgtacg taagctcggg attcggctcg agcttgtctt ggaccttgaa 60  
 tctcacaagt tgcaagcttt cactgangca tatgcagaac ttgcacctgc tttgtctgat 120  
 ctgaatgttc ttgttgagac ctactttgct gacatccctg ctgaggcgta caagaccctc 180  
 acatctctga atggcgtcac tgcataatggg tttgatttgg tccgtggaac ccatactctt 240  
 gatttgatca aggttggtatt tcccagtgga 270

<210> 2952

<211> 549  
 <212> nucleic acid  
 <213> Glycine max  
  
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 tcneggggtcg acccacgcgt ccggcagatc tgagaccacc taccagattg ctttgtctat 120  
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 tggatcatcg canatcactg aattanaaat tttttttggt natcctnatt ttcacatatg 360  
 tttgggnataa ncaantttnc gtatngacag atccannact canatgtgnc tactcggact 420  
 tcaanegact ntntccaat tncattannt nancntggan tgcntgangt ntatgnnonn 480  
 nttnnnannt ttntgtngna tganaagtag gttntnttn atngntatag tnnnanggtt 540  
 ttnttgtn 549

<210> 2953  
 <211> 317  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2953  
  
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 agttccaatc ctgctcaacc accacnnttg gatccttccc tacagactgt agaactgagg 120  
 anggtacgcc gtgaattcna ggctaacaag atctccgagg aagagtatgt naagtcaatt 180  
 aaggaggaaa ttcgcaaagt tgttgagctt caagaagagc ttgatattga tgttcttgtt 240  
 catggagaac cagagagaaa tgatatgggt gagtnttcgg tgaacaattg tcaggcttgc 300  
 ttcaccgtta atgggtg 317

<210> 2954  
 <211> 321  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2954



anatgcang cacgntacg taagctcgga attcggtcg agttgngcnc tgntgtggac 60  
 ttcacctatg ctttctcana aggtgtnga tgaatacnag gngggccaag ggcgcttnga 120  
 gtggatacng ttccggnect cgttggcnct gttacatagc tgttgccttc caagcctgcc 180  
 aagggagttg ngaaatcctn ttctctcctc tctctcctc ncaaggttct tgctgtctac 240  
 aaggaagtta ttgntgacct taaggcagct ggtgcttcat ggattcagtt tgatgagcct 300  
 acccttggtc ttggaccttg n 321

<210> 2955  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max

<400> 2955

gtgcngcan gngtacgtnn agctcggaat tcggctcgag gtgagggtag gccgtgaatt 60  
 caaggctaac aagatctccg aggaagagta tgtaaagtca attaaggag gaaattcgca 120  
 aagttgttga gcttcaagaa gagcttgata ttgatgttct gttcatggag aaccagagag 180  
 aaatgatatg gttgagtact cgggaacaa ttgtcaggct tgctcaccg ttaatgggtg 240  
 ggtgcaatcc tatggttccc gttgcgtgaa gccaccgatc atctatggtg atgtgagccg 300  
 cccaaagcca tgaccgtc 318

<210> 2956  
 <211> 260  
 <212> nucleic acid  
 <213> Glycine max

<400> 2956

ncgcatgcac gcgtacgtna gctcggaatt cggctcgagg gaggccaagg cgcttgagat 60  
 ggataccgtt ccggtcctcg ttggccctgt tacatacctg ttgctctcca agcctgccaa 120  
 gggagttgag aaatcctttt ctctcctctc tctccttccc aaggtcttgc tgtctacaag 180  
 gaagttattg ctgaccttaa ggcagctggt gcttccatgg attcagttgg nggagctaac 240  
 cctggtctgg gacctgnngt 260

<210> 2957  
 <211> 247  
 <212> nucleic acid

<213> Glycine max

<400> 2957

cgtcgcangc acgcgtacgt aagctcggaa ttccggctcga gctgccactg aggaaatcag 60  
aacaagctca ctaccttgga ctgggctgtc catgccttca gaatcnnona tnttgngng 120  
cangatacna ctcagatcca caccacatg tgctactcca acttcaacga catcatccac 180  
tccatcatcg acatggacgc tgatgttatc accattgaga actctcgctc cgntgagaa 240  
gctcctg 247

<210> 2958

<211> 187

<212> nucleic acid

<213> Glycine max

<400> 2958

gggaatgctt ancggtcntg ttancantct caannngtcc tttgttagaa atgaccaacc 60  
tagatctgag accantacc agattgcttt gtctatcaag gacgangtgg aagacntga 120  
aaaggctggc atcantgtna tccaaattga tgaagctgct tggagagagg gttncaccagt 180  
gangaat 187

<210> 2959

<211> 250

<212> nucleic acid

<213> Glycine max

<400> 2959

aataccacca actgaagaaa ttgctgacag aatcaacaag atgctggcag tgctcgagaa 60  
gaacatcttg tgggtgaacc ctgactgtgg gctcaagacc cgtaagtaca ctgaggtgaa 120  
gccagccctc acaaacatgg ttccgcagc aaaactcatc cgcaacgaac ttgccaagt 180  
aatgggtata ggaagtngan tttccaagtn atgggggtccg ntttaattta aaaccccccc 240  
aaaaaaattt 250

<210> 2960

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2960

gtcgcangca cgcgtacgta agctcgggaat tcggctcgan ctcgagccga atcggctcga 60  
gataccacca actgaagaaa ttgccgacag aatcaacaag angtcggcag tgctcgagaa 120  
gaacatcttg tgggtgaacc ctgactgttg gctcaagacc cgtaagtaca ctgaggtgaa 180  
gccagccctc acaaacatgg ttgccgcagc aaaactcatc cgcaacgaac ttgccaagtg 240  
aatggtataa gaaagtagaa tcttccaagt catttggttc tgctttatat tat 293

<210> 2961

<211> 261

<212> nucleic acid

<213> Glycine max

<400> 2961

cnagnattca ctgacgcata tgnagaactt gcgccacgct ttgtctgggt tgacatgttc 60  
ttnttgagac ctactttggc tgacatccct gctgaggcat acaagaccct cacatctctg 120  
aatggcgtca ctgcatatgg atttgatttg gtccgtggaa ccaanactct tgatttgatc 180  
aaggggtggat ttcccanggg aaaatacttt tttgggggan tgntgatgga aggancattg 240  
ggccaatgac tttgctgttt t 261

<210> 2962

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2962

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tcctttctct cctctntntn cttcgnaagg ntnttgngt ctanaaggaa gtnattgntg 180  
accttaaggc agctgggtgc tcatggattc agtttgatga gcctaccctt gtcttgacc 240  
ttgngtctca caagttgcaa gcattcagtg ccgcana 277

<210> 2963

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2963

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tgatattgat gttcttgttc atggagaacc agagagaaat gatatggttg agtacttcgg 180  
tgagcaattn tnaggctttg cttcactgt taatggntgg gtgcantcca tggttcccgt 240  
tgtgtgaagc aacaatnnac caaggnnatt aaccgccca aagccattga ott 293

<210> 2964

<211> 227

<212> nucleic acid

<213> Glycine max

<400> 2964

accaactgaa gaaattgctg acagaatcaa caagatgctg gcagtgctcg agaagaacat 60  
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cctcacaaac atgggtgccg cagcaaaact catccgcaac gaacttgcca agtgaatggg 180  
ataagaaagt agaatcnnac caagtcattt gggtctgctt tatatta 227

<210> 2965

<211> 290

<212> nucleic acid

<213> Glycine max

<400> 2965

ntcgangca cgcgtacgta agctcggaat tcggctcgag gtgaccaacg aggctgttca 60  
gaaggntgct gctgcattga agggtncaga tcatcgccgt gcaacaaatg tcagtgccag 120  
actggattct caacaaaaga agctcaacct tccaatcctg nccaaccacc actatnggat 180  
ccttcctca gactgtagaa ctgaggaggg naggcngngaa ttcaaggcta acaagatctc 240  
cgaggaagag tatgtaaaagt caattaagga ggaaattcgc aaagttgttg 290

<210> 2966

<211> 256

<212> nucleic acid

<213> Glycine max

<400> 2966

aagacctcac atctctgant gtnctcant gcatatgggt ttgatttggt cegtgaago 60  
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 ttgatggaan gaacatcngg gccaatgacc ttgtgtcttc tcncaactaca tgcaggggtct 180  
 tgagggcatt gtgggcaaag ataagcttgt tgtgnccacc tcnncctccc ttctcacact 240  
 gctgtngatc ntgtna 256

<210> 2967  
 <211> 330  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2967

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 agaatcaaca agatgcacgc agtgctcgag aagaacatct tgtgggtgaa ccctgactgt 180  
 gggctcaaga cccgtaagta cactgagggtg aagccagccc tcacaaacat ggttgccgca 240  
 gcaaaaactca tccgcaacga acttgccaag tgaatgggtat aagaaagtag aatcttccaa 300  
 gtcatttggt tctgtttata ttataataca 330

<210> 2968  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2968

tcgcangcac gcgtacgtaa gctcgggttc gacaccanct acnacatnaa ttgtccctga 60  
 attgggccct gatgtgaact tcacctatgn cnotcacaag gctgttgatg aatacaagga 120  
 ggccaaggcg cttggagtgg ataccgttcc ggtcctcgnt ggccctgtta catacctgtt 180  
 gctctccaag cctgccaagg gagtnagaaa gccntttcgc tcnctctct cgggcccaag 240  
 gacttgctgt cnacaaggaa gnnattgcng accngaange agcnggngca tcanggatca 300  
 gttnga 306

<210> 2969  
 <211> 215  
 <212> nucleic acid

<213> Glycine max

<400> 2969

tnagctcgga attcggtctg agcanggcgt tggagtggat accgttcggg tctctgttgg 60  
ccctgttaca tacctgttgc tcttccaaag cctgccaaag gagttgagaa atccttttct 120  
ctcctctctc tcttcccaa ggttcttgcgt gtctacaagg aagttattgc tgaccttaag 180  
gcagctggtg cttcatggat tcagtttgat gagcc 215

<210> 2970

<211> 172

<212> nucleic acid

<213> Glycine max

<400> 2970

ngtcgcatgc acgcgtacgt aagctcggaa ttcggtctga gcgcatatgc agaacttgcg 60  
cctgctttgt ctggtttgaa tgttcttgtt gagacctact ttgctgacat ccctgctgag 120  
gcatacaaga ccctcacatc tctgaatggc gtcactgcat atggatttga tt 172

<210> 2971

<211> 170

<212> nucleic acid

<213> Glycine max

<400> 2971

gtcgcangca cgcgtacgta agctcggaa ttcggtctgag ngcttctcac aaggctgttg 60  
atgaatacaa ggaggccaag gcgcttggag tggataccgt tccggtcctc gttggccctg 120  
ttacatacct gttgctctcc aagcctgcc aaggagttga gaaatccttt 170

<210> 2972

<211> 321

<212> nucleic acid

<213> Glycine max

<400> 2972

gtcgcangcn cgcgtacgtn agctcgtcaa ttcggtctga gnacttgtn gctctccaag 60  
cctgnccaag ggagtcgaga aatccctttt ctctcctctc tctccttccc aaggttcttg 120  
ctgtctacaa ggaagttatt gctgacctta aggcagntgg tgcttcatgg attcaatttg 180

atgagcctac cctgtctgga ccttgaatct cacaagttgc aagctttcac tgacgcatat 240  
gcagaacttg gcacctgctt tgtctgatct gaatgttctt gtngagacct atcntgctga 300  
catccctggt gngnggtana a 321

<210> 2973  
<211> 236  
<212> nucleic acid  
<213> Glycine max

<400> 2973

tacaaggagg ccaaggcgct tggacgtgga taccgttccg gtccctgcgtt ggccctgtta 60  
aatacctggt gctctccang cctgncaang gagttgagaa atccttnnct ctccctctctc 120  
tccttcccaa ggttcttgct gtctacaagg aagttattgc tgaccttaag gcagtgggtgc 180  
ttcatggatt cannttnatg agtctacnct gtnttggact tgagtctcac aagttg 236

<210> 2974  
<211> 231  
<212> nucleic acid  
<213> Glycine max

<400> 2974

actcnanncn cncntncgtn agcncgggnt tcggctcnag nttggntcnt tcnctcagac 60  
tgtanaacng ngnagggtac gccgtnaatt caaggctaac aanatctgcn gnggangagt 120  
atntaaagtc aattanggag gaaattcgca aagttgttga gcttcaagaa gagcttgata 180  
ttgatgttct tgttcatgga gaaccagcga nanatgntat ggttnagtcc c 231

<210> 2975  
<211> 313  
<212> nucleic acid  
<213> Glycine max

<400> 2975

tnntngcacg cgtacgtaag ctcggaattc ggctcgagat gttgtaatat ttattctgct 60  
gttactcatg gcttcttttc ttctctctca ggctgctgca ttgaagggtt cagatcatcg 120  
ccgtgcaaca aatgtcagtg ccagactgga ttctcaacaa aagaagctca accttccaat 180  
cctgnccaac caccactatt ggatccttcc ctccagactgt agaactgagg agggtagcgc 240

ggaattcaag gctaacaaga tctccgagga agagtatgta aagtcaatta aggaggnnat 300  
tcgcaaagtt gtt 313

<210> 2976  
<211> 184  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2976

atncagcgta cgtaagtcgg aattcggtc gagcagagtt taccaagcgc ccaatgaagg 60  
gaatgcttac cggctctggt ancattcttc aactggctct ttgtagaaa tgaccaacct 120  
agatctgaga nccantacca gattgctttn ggctatcaaa gacgaantng agggnncttg 180  
aaaa 184

2977  
314  
nucleic acid  
Glycine max  
2977

<210> 2977  
<211> 314  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2977  
  
nngtngcang cacgcgtacg taagctcgga attcggtctg aggggtgaacc ctgactgtgg 60  
gctcaagacc cgtaagtaca ctgaggtgaa gccagccctc acaaacatgg ttgccgcagc 120  
aaaactcatc cgcaacgaac ttgccaagtg aatgggtataa ganagtagaa tcttccaagt 180  
catttggttc tgctttatat tataatacac caaagaaaaa ttttctctat attgggttgt 240  
ttcaataact gtgtgtggaa tatttaggtg tcttagcatg ctctgtgagc aattgattct 300  
tcctcaaccc ctcc 314

<210> 2978  
<211> 153  
<212> nucleic acid  
<213> Glycine max  
  
<400> 2978

ncgcatgcac gcgtacgtaa gctcggaatt cggctcgagg tgaccaacga ggctgttcag 60  
aaggctgctg ctgcattgaa gggttcagat catcgccgtg caacaaatgt cagtgccaga 120  
ctggatttct caacaaaaga agctcaacct tcc 153



<210> 2979  
 <211> 280  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2979  
  
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 gctgggtgctt catggattca gtttgangag cctnnncttn tnttggaccc ngngnanaa 120  
 ngnnnganga nnncgcnag gaanatggga attntgcgcg tacttngnnn ngmntgaang 180  
 annntnatna naccctctnt nntggaatac cnnatngnngn aaccnngaac cctnnecatct 240  
 ctggaatggc gnnatgcgta tggattgatn agtcngtgga 280

<210> 2980  
 <211> 102  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2980  
  
 cccccagtgg anaatacctc nttgctggag tggttgatgg aaggaacatc tgggccaatg 60  
 accttgctgc ttctctccac tacattgcag ggtcttgagg gc 102

<210> 2981  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2981  
  
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 cccgtaagta cactgaggtg aagccagccc tcacaaacat ggttgccgca gcaaaaactca 120  
 tccgcaacga acttgccaag tgaatggat aagaaagtag aatcttccaa gtcatttggt 180  
 tctgctttat attataatac accaaagaaa aattttcccc atattgggtg nttnataac 240  
 tgnngtgga atatttangt gncttagcat gctctgtgag caattgat 288

<210> 2982  
 <211> 260  
 <212> nucleic acid  
 <213> Glycine max

<400> 2982

agctcggaat tcggctcgag cgtaagtaca ctgaggtgaa gccagccctc acaaacatgg 60  
ttgccgcagc aaaactcctc cgcaacgaac ttgccaaagt aatggtatga gaaagtagaa 120  
tcttccaagt catttggttc tgctttatat tataatacac aaagaaaaat tttctctata 180  
ttgggttggt tcaataactg tgtgtggaat atttaggtgt cttagcatgc tctgtgagca 240  
attgattctt cctcaacccc 260

<210> 2983

<211> 323

<212> nucleic acid

<213> Glycine max

<400> 2983

gtcgngcac gcgtacgtaa gctcggaatt cngctcgagc totgcaactct ctctttctca 60  
tcctcttctc ttctgttctc tgttcgtgcc acttcttctc cgagcaatgg gcatctcata 120  
ttgttggtta tccacgcatg ggaccaaga gagaacttaa gtttgctttg gaatcttttt 180  
gggatggaaa gagtagtgct gatgatctgc agaaggttgc tgetgacctt aggnacagcca 240  
tctggaagca gatggctgat gctggaataa agtatattcc tagcaacact ttctcatact 300  
atgatcaagt actggacaca acn 323

<210> 2984

<211> 335

<212> nucleic acid

<213> Glycine max

<400> 2984

ngtcgcangc acgctacgt nagctcgga ttcggctcgn nctcgtctctg cactctctct 60  
ttctcatcct attctcttcg cttctctggt cgtgccactt cttctnagag caatggcatc 120  
tcatattggt ggatatccac gcatgggacc caagagagaa ctttaagtttg ctttggaatc 180  
tttttgggat ggaaagagta gtgctgagga gctgcagaag gttgctgcag accttaggtc 240  
agccatctgg aagcagatgg ctgatgctgg aataaagnat attcctagca acaccttctc 300  
actttacgat caagtatgga cacaacagcc atgct 335

<210> 2985

<211> 297  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2985  
  
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 actctctcct tctcatcctc ttctcttcgc ttctctgttc gtgccacttc ttctcgagca 120  
 atggcatctc atattgttgg ttatccacgc atgggaccca anagagaact taagtttgct 180  
 ttggaatctt tttgggatgg aaagagtagt gctgatgac tgcagaaggt tgctgctgac 240  
 cttagggtcag ccatctggaa gcagatggct gatgctggaa taaagtatat tcctagc 297

<210> 2986  
 <211> 327  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2986  
  
 nnttancaag cgtangtaag ctcggaattc ngctcgagct cactctgcac tctctcctct 60  
 ctcatectcg tgtctnccn ncnntcggnc ngcnattccg tcccagacna tggcatctca 120  
 tattgtnggt tatccacgca tggccnccca ngagaganct taagttnngct ttggaatctt 180  
 tttgggatgg aaagagtagt gctgatgac tgcnagaaggt tgctgctgac cttagggtcag 240  
 ccatctggaa gcagatggct gatgctggaa taaagtatat tcctagcaac actttctcat 300  
 actatgatca agtactggac acaacag 327

<210> 2987  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 2987  
  
 cnatcgcang cangcgtacg taagctcgga attcggctcg aggtgaagtg antagtataa 60  
 atntacaaan tcncctncan tcattgtgtg gcgccgttna cattctcgtnt ctgcactctc 120  
 tctttctcat cctnntctct tcgcntctct gtncgtgccca cntcttctcg agcaatggca 180  
 tctcatattg ttggttatch acgcatggna cccaagagag aacgtaagtt ngcttnggan 240  
 tctntttggg atggaaagag nagtgctgag gagctgcaga aggttgctgc agaccttagg 300

tgagccatct tgaan 315

<210> 2988  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2988

agtcgcatgc acgcgtacgt aagctcggaa ttcggctcga ggattattaa tcgtatcagt 60  
 gaaagaagta agaagagaga gaagtgaagt tagtagtata aatgtacaaa ctctcctcta 120  
 ttcagtgtgt ggcgccgttt acgatctcac tctgcaactct ctctcttctca tctcttcttc 180  
 ttcgcttctc tgttcgtgcc acttcttctc gagcaatggc atctcatatt gttggttatc 240  
 cacgcatggg acccaagaga gaacttaagt ttgctttgga atcttttttg gatggaaaga 300  
 gtagtg 306

<210> 2989  
 <211> 264  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2989

gtcgcangca cgcntacgtn agctcggaat tcggctcgag ctctgcactc tctctttcnc 60  
 atcctattct cttcgcttct ctgtncogtg ccacttcttc tcgagcaatg gcatctcata 120  
 ttgttggtta tccacgcatg ggaccaana gagaacttaa gtttgctttg gaatcttttg 180  
 gatggaaaga gtagtgctga ggagctgcag aaggttgctg cagaccttag gtcagccatc 240  
 tggaagcaga tggctgatgc tggc 264

<210> 2990  
 <211> 316  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2990

tacgcgtacg taagctcgga attcggtcgc agcttctttc cctaattgac aagatccttc 60  
 ctgtctacag ggaggntggt gctgaattga aggcagctgg tgctacttgg atccagtttg 120  
 atgaacctac ccttgtgaag gatctcaana cccaccagtt acaagcattt acacatgnct 180

atgcagagct agagtcaagt ttatctggtt ttaatgttct gattgagana tactttgctg 240  
atgtccctgc tgaagcatac aaaacactca cctctttgaa ggctgttact gcatatgggt 300  
ttgatattgt tcgtgg 316

<210> 2991  
<211> 321  
<212> nucleic acid  
<213> Glycine max

<400> 2991

caanatacat gcacgcntac gtnagctcgg aattcggctc gagggcagct ggtgctactt 60  
ggatccagtt tgatgaacct acccttgtga aggatctcaa caccaccag ttacaagcat 120  
ttacacatgc ctatgcagag ctagagtcaa gtttatctgg ttttaatgtt ctgattgaga 180  
catactttgc tgatgtccct gctgaagcat acaaaacact cacctctttg aaggctgtta 240  
ctgcatatgg gtttgatatt gttcgtggaa caaagaccct ggatttggtc naggcaggtt 300  
ttcccneggg gaaatntttt t 321

<210> 2992  
<211> 331  
<212> nucleic acid  
<213> Glycine max

<400> 2992

ttgcangcac gcgtagtaag ctcggaattc ggctcgagnn tctttcccta attgacaaga 60  
tccttcctgg tctacagga ggtngttgct gaattgaagg cagctggtgc tatttggtac 120  
tagtntgatg aacctacnt tgtgaaggat ctcaacaccc accagttaca agcatttaca 180  
catgcctatg caganctaga gtcaagttta tctggtttta atgttctgat tgagacatac 240  
tttgctgatg tcctgtctga agcatacaaa aactcacct ctttgaaggc tgttactgca 300  
tatgggtttg atattgttcg tggaacaaag a 331

<210> 2993  
<211> 284  
<212> nucleic acid  
<213> Glycine max

<400> 2993

gatcgacatg cttgggctgg agaatacccc ggcgtgaagc gcatcaccat caagcccca 60  
 actgacagat ggggtctccct gagaccaaca cggatatcatg tcttggctga gggtcgattg 120  
 atgaacttgg atgcgcaatg gaaaccccag tttggagtct gttcctcaac naccagtcac 180  
 tgtcacttag tgtgaagggg natacgnagt acgagagagt tagttgccca gactgagaga 240  
 gtgtgcntta ctggcaattg gactagtgcc actacagccg gtgt 284

<210> 2994  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2994

cangcacgcg taogtaagct cggnatctcg ctcgaggggc cccgnacggc ggccccgacc 60  
 tnatcgctga cgacggagge gacgccacc tctcatcca cgagggcgte aaggccgagg 120  
 agctctatga gaagaccggg gaactcccc accctaact cactnanaat ncnanmntc 180  
 cagatcgctg ttaccnncan cagnganngg ttgaanaccg atcncaccan gtaccgnaag 240  
 atgaaggncg gtctcgcttg ggtttctgag aaaccaacac tgggtgtaag agnctat 297

<210> 2995  
 <211> 318  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 2995

ngtcgcatgc acgcntacgt aagctcggaa ttcggctcga gtgagaagac cggggaactc 60  
 cccgacccta actccactga caacgccgag ttccagatcg tgcttaccat catcagagat 120  
 gggttgaaga ccgatccac caggtaccgc aagatgaagg agcgtctcgt tggggtttct 180  
 gaggaanaac aaccactggt gttaaagang gctaaatccg gatgncaggn gaatggggat 240  
 tctantcntt ccctgnngat aaatgtcaat gagannctnt tcnnaaggn ccangtttgn 300  
 aaanttttna cgggnccg 318

<210> 2996  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max

<400> 2996

cgtcgcangc acgcgtacgt aagctcggaa ttgggtcga gcccgacccc anctccaccg 60  
ncaacgccga gtttcanatc gtgcttacna tcatcagaga tgggttgang accgatccca 120  
ccaggtaccg caagatgncg gngcgtctcg ttgggggttc tgaggnnacc accactggng 180  
tnangaggct ctatnagatg naggcgaatg ggncctctct ctccctgcn attnntgtna 240  
atgacnngtc nccangagca agtttgacaa cntgtatngg tgccgtca 288

<210> 2997

<211> 334

<212> nucleic acid

<213> Glycine max

<400> 2997

cgtcgcngca cgogtacgtn agctcggaa tgggtcga tcaacatcaa gcctcaaaca 60  
gacagatggg ttttccccga taccaagagg gggatcatcg tgttggcaga gggtcgtttg 120  
atgaacttgg ggtgtgccac gggacacccc agctttgtga tgtcgtgctc cttcaccaac 180  
caggtcattg ctcagcttga attgtggaaa gagaagggtt ctgggaagta tgagaagaag 240  
gataatnnat nccaagcac ctnacgagaa agnngcntna nanctaccat tgccagcatt 300  
gagcnntgcg naccaagcgt tccaaagacc aagc 334

<210> 2998

<211> 277

<212> nucleic acid

<213> Glycine max

<400> 2998

gcacgcgtac gtaagctcgg aattcggctc gagcggagtt ccagatcgtg ctgagcatca 60  
tcagggatgg cttgaagacc gatcccaaga ggtaccacaa gatgaaggac agaatcgtcg 120  
gtgtctccga agaaaccacc accggtgtca agaggctcta ccagatgcag gccaatggct 180  
ccctcttgtt ccctgccatc aacgtcaatg actcggtcac caagagcaag tntgataact 240  
tgtatggatg ccgtcactcg cttcccgatg gactgat 277

<210> 2999

<211> 293

<212> nucleic acid

<213> Glycine max

<400> 2999

tcgcangcac gcgtacgtaa gctcgnaat tcggtcgcag ctcagccctt caagggggcc 60  
cgcatcaccg gctcccttca catgaccatc cagaccgtg tctcatcga gaccnncanc 120  
gttctngggc ncgtgggttc ntngtgcctc tnnaanntnt tcnccattna gnnnncanc 180  
gengtcgna tcgncgtnt cagcgtccgc tgtctacgcc tgtaatggcg ataccotcca 240  
tgagtactnt tgggtgcacng atnggccctc gatgggnccc cggcgngcn ccc 293

<210> 3000

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 3000

gcgaangcat gcacgcgtac gtaagctcgg aattcggctc gaggccagct tggagctagg 60  
ctcaccaagc tttccaaaga ccaagctgat tacatcagtg tgctgttga gggccatac 120  
aagcctctc actacaggtc ctgatccatc ctattnnggg agaataaacc taaactattn 180  
tatcaattcc cgaggcntca ttgttacact ttcctttttg gattttttcc attacaattt 240  
acntttgtgg tagcatcgga gcttcttttt tcttttttag tannatca 288

<210> 3001

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 3001

gcangcacgc gtacgtaagc tcggaattcg gctcgaggtc tccgaggccg acatcttcgt 60  
caccaccaca gggaacaagg acattatcat gcttgaccac atgaagaaga tgaagaacaa 120  
tgccatcgtg tgcaacattg gccacttcga caacgaaatc gacatgctgg ggcttgagac 180  
ctgccctggg gtgaagcgca tcaacatcaa gcctcanacn gacagatggg ttttccccga 240  
taccaagagg gggatcatcg tgttggcaga gggctgtttg atgaan 286

<210> 3002

<211> 294

<212> nucleic acid



<213> Glycine max

<400> 3002

acgncgcang gacncgtaag ntcagggtctc nagctcgngg cnaggatccc aagaggtacc 60  
acaacgnnga aggacagaat cgtcgggtgtc tccgaagana ccaccaccgg tgtcaagagg 120  
ctctaccaga tgcnggcnaa tggctccctc ttgttccttg ccatcaacgt caatgactcg 180  
gtcaccaana gnangtttga taacttgtnt ggatnccgtc actcgcttcc cgatggactg 240  
atgagagcca ctgatntgat gattgccgga aaggngactg ttgtgtgtgg ctac 294

<210> 3003

<211> 256

<212> nucleic acid

<213> Glycine max

<400> 3003

gtcacacgag gtcccgatag antgatgaan cgccactgat gtgatgatng ccggaaaggt 60  
agcagttgng gnnggcaccg agangnggca aggntagnnc ancgcnana ngnaaaccng 120  
ggcncgtgtc atngtcancg agatngatcc catntntncc cttnaggcnc taatgncggg 180  
tcttcaggtt ctcacctgg aagatgtngt ctccgaggcc gacatntcg tcaccancac 240  
agggaacaag gacatt 256

<210> 3004

<211> 302

<212> nucleic acid

<213> Glycine max

<400> 3004

gtngcannca cgcgtacgta agctcggaat tcggctcgaa ngggccccgg cggcggcccc 60  
gacctcatcg tcgacgacgg aggcgacgcc anctcctca tccacgaggg cgtcaaggcc 120  
gaggagctct atgagnagac cggggaantc cncgnnccta antccactna caacgccgag 180  
ttccagattg tgcttaccat catcagngat gggttnnaana angatccan naggtaccgn 240  
aanntgaagg ancgctctcg tggggtttct gangaaccga cgatgggtgtt aagatgtana 300  
nc 302

<210> 3005

<211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3005  
  
 aaanontcaa tgcangcac gcgtacgtna gtcggaatt cggctcgagc agagatgggt 60  
 tgaagaccga toccaccagg taccgcaaga tgaaggagcg tctcgttggg gttnctgagg 120  
 aaaccaccac tgggtgtaag aggctatata agatgcagcg attgggatcc tatntttccn 180  
 ngnaattaat gtcaatgact ctgttaccaa gancaatttg acaacttgta ccgggtgccg 240  
 tcantctctc cctgatggnc tantnaggcc tactnntgtg ntgttgncgt gaaagttggn 300  
 ngtgttgcnt gat 313

<210> 3006  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3006  
  
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 ctccacatga cnatccagag gagcggttnt cattgagann ntcanngncc ttggngncga 120  
 ggtncgntgg tgctcntgca anatcttctc caccaggan cagcnnacg cngctattgc 180  
 ccgcanagt gcngccgtct tcgntggaa gggtagagacc ctccaggagt actggtggtg 240  
 caccgagcgc gcnetcgant ggggccccgg tggtaggaccg anctcatcgt cgacgaggtg 300  
 gtgang 306

<210> 3007  
 <211> 70  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3007  
  
 attnttntgc ttgaccaca innagaanat gaagnannan tgccttggtg ttgcaacatt 60  
 tggccacttt 70

<210> 3008  
 <211> 536  
 <212> nucleic acid

<213> Glycine max

<400> 3008

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cgcgctccgcc cacgcgtccg nccgntgcga naagannaca gaagggggcc caggctgatt 120  
acatcantgt gcctgttgag ggtccatnca agcctgctca ntacnngtac taagtaattg 180  
agattatcaa cggaacagtg aggganagac ntantcgggtt ttatgaatcg ggntgattgt 240  
ttaagtnttc cttttttttg aggttttgtt gttanacttt tcagatttga gggtagcctn 300  
agtttanctt tngggcngcn naagnagnag tcaggtnttn aaaaaaggng gcngngntgg 360  
nngatcnaa nntaacgtac cttgcntnca acgtcatnnc tcttcgaaag tggcaccnat 420  
tttcaattca ggggcgggnc gttttaannn cnnctttnnc ggggaaaacc ttggggntan 480  
ccanggttac ccccttgnen tnanncccn ttttcccnna nttgggttaa aaaaaa 536

<210> 3009

<211> 330

<212> nucleic acid

<213> Glycine max

<400> 3009

gtccnatncc gctgngngna gacnggggt tgggtnacnt ggnntcntc ntggttgncc 60  
ncttgagggn cttgatganc aatgccattg tttgctncat tggtcacttt gggcctgaga 120  
tngacntgct tggntcggag nactattccg gtgtgntnng catcaccatc atgncccctt 180  
tctgacagat gggctctccc tgataccngc nccggtatcn ttgtcttggc tgagggtcnn 240  
ttgntnatct tgggatncnc cattnngnac ttcctttttg tgatgtcctg ctccttcngc 300  
antngntcn ntgctcngnt tgngttggtg 330

<210> 3010

<211> 473

<212> nucleic acid

<213> Glycine max

<400> 3010

gtttgattcc gagccannca atgcatcgna nncangcgta cntaaactcg gaattcggcn 60  
cnagcaatga ctctgtcacc aagagcaant ttgacanctt gtatgggtgc cgtcactctc 120

tccctgatgg notcatgagg gctaccgatg ttatgattgc tggaaagggtg gctgttgtgg 180  
ctggatatgg tgatgttggc aagggttgtg ctgctgcaat gaagcaggct ggtgctcgtg 240  
tcacgtgac cganattgat cccatctgtg cccttcaggc tctcatggaa ggccttcagg 300  
ttctgacctt ggaggatgtt gtttctgagg ctgatatctt tgtcaccacc accggttaaca 360  
aggacatcat catggttgac cacatgagga aaatgaacaa caatgccatt gtttgcaaca 420  
ttggtcacnt tgacaatgag atcgacatgc ttgggctgga gaactaaccg ggg 473

<210> 3011  
<211> 500  
<212> nucleic acid  
<213> Glycine max

<400> 3011

gaggnntgnt ttgnanchnan anactttcgc ctgccgtacc ggtccggaat tcccgggtcg 60  
acccacgcgt ccgcggacgc gtgggcggac gcgtgggctg cgagagacga cagaaggggg 120  
gactctactc ttccctgcta ttaatgtcaa tgactctgtt accaagagca agtttgacaa 180  
cttgtagcgg tgctgtcact ctctccctga tggctgatg agggctactg atgtgatgat 240  
tgctggaaaag gtggctgttg tggctggata tggngatgtt ggcaagggtt gtgctgctgc 300  
attgaagcag gctgggtgctc gtgtcatcgt gactgagatt gaccccatth gtgcccttca 360  
ggctctcatg gaaggccttc aggttctgac cttggaggat gttgtttctg aggctgatat 420  
ctttgtcacc accacgggta acaaggacat catcatggnt gaccacatga agaaaatgaa 480  
gaacaatgcc attgtttgca 500

<210> 3012  
<211> 383  
<212> nucleic acid  
<213> Glycine max

<400> 3012

ccattgtttg caacattggt cactttgaca atgagatcga catgcttggg ctggagaact 60  
acccoggcgt gaagcgcac accatcaagc cccaaactga cagatgggtc ttccctgaga 120  
ccaacaccgg tatcattgtc ttggctgagg gtcgattgat gaacttggga tgcgccactg 180  
gacaccccag ttttgtgatg tcctgctcct tcaccaacca ggtcattgct cagcttgagt 240

tgtggaagga gaagagtacc ggcaagtacg agaagaaggt ttacgttttg cccaagcacc 300  
 ttgatgagaa ggtggctgca cttcacctgg gcaaacttgg agctaagctg acccagctta 360  
 gcaagtccca ggctgattac atc 383

<210> 3013  
 <211> 528  
 <212> nucleic acid  
 <213> Glycine max

<400> 3013

gnnnggaggt ttganngggg gngggnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60  
 nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnagagagan agagagagat ctatctatct 120  
 atcaagatgg ngttgttggg tgagaaaaca agcagtggaa gggagtacaa ggtgaaggac 180  
 atgacgcaag ccgatttcgg aagattggaa atcgagctgg cggagggtga aatgcccggc 240  
 ctcatgtect cccgcaccga gttcggcccc tctcaacct tcaagggcgc taggatcacc 300  
 ggctccctcc acatgaccat ccaaaccgcc gtccctcatg agaccctcac cgccctcggc 360  
 gccgaggtcc gctggtgctc ctgcaacatc ttctccaccc aggaccatgc cgccgccgcc 420  
 atngcccgcg acagcgcctt cgtcttcgcc tgggaaggggt gagaccctnc aggaatactg 480  
 gtggtgcacc gagcgcgcc tgcactgggg ccccggcggn gggcccca 528

<210> 3014  
 <211> 520  
 <212> nucleic acid  
 <213> Glycine max

<400> 3014

agngtttttg tanntggggg gggggggggn aagnganata tnttagctat agatnlnaca 60  
 tgtacanngt acgtaagctc ggaattcggc tcgagggccg acttcggccg cctcgagatc 120  
 gagctggccc gaggttgaga tgccgggcct catggcctgg cggaccgagt tcggcccatc 180  
 tcacccttca agggggcccg catcaccggc tcccttcaca tgaccatnca gaccgctgtc 240  
 ctcatcgaga cctcaccgn tctcggcgcc gaggttcgct ggtgctcctg caacatcttc 300  
 tcaactcagga ccacgccgcc gncgccatcg cccgtgacag cgccgncgtc ttcgcctgga 360  
 agggtgagac cctccaggag tactggtggt gcaccgagcg cgcccttgac tggggccccc 420

gcggcgggccc cgacctnatt gtcgacgacn gaagcnacgc cacctctnat cacgaaggcc 480  
tnaangncca gggctctatn annaagaccg gggaactccc 520

<210> 3015  
<211> 344  
<212> nucleic acid  
<213> Glycine max  
<400> 3015

gttattttctc agcgcgtaaa gcatggcttt gttggtggag aaaaccacga gtggtcgcga 60  
gtacaaggtc aaggaccttt cccaggccga cttcggccgc ctcgagatcg agctggccga 120  
ggttgagatg cccggcctca tggcctgtcg naccgagttc ggcccntccc agccttcaag 180  
ggggcccga tcaccggctc cotccacatg accatccage ggcgccgttc tcattgagac 240  
cctcaccgcc cttggcgccg aggtccgctg gtgctcctgc aacatcttct ccanccagga 300  
ccacgccgcc gccgtanttc ncgcgacagt gngccgtct tcgc 344

<210> 3016  
<211> 528  
<212> nucleic acid  
<213> Glycine max  
<400> 3016

ggnnnaagtt tttgnngggg ggggnannan gtnnnnnnnnn nnnnnnnnnn nnnnnnnnnn 60  
nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnnnnnnnnn nnntcgagat cganctggcc 120  
cgaggtnag atggcgatcc tcatggcggtg gcggaccgag ttcggcccat ctcancctt 180  
gaagggggcc cgcathaccg gctccnttca catgaccatt canaccgntg tctcatoga 240  
naccctnacc gctctcggcg ccgaggttcg ctggtgctcc tgcaacatct tctccactca 300  
ggaccacgcc gncgncgcc tgcgccgtga cagcgccgnc gtcttcgcct ggaagggtga 360  
gaccctncan gactactggt ggtncaccga gcgcgccctt gactggggcc cccgcgnggc 420  
cccgacctta tngtcnaccg accgaaggca accccacctt cttatcaacn aggcgtnaa 480  
ggccaaggag ctctatnaag aagaccgggg aactccccna cctaact 528

<210> 3017  
<211> 347  
<212> nucleic acid

<213> Glycine max

<400> 3017

cgggtccct tcacatgacc atccagaccg ctgtcctcat cgagaccctn caccgctctc 60  
ggcgccgagg ttcgctggtg ctctgcaac atctttctcca ctcaggacca nnnnnnnnnn 120  
nnnnnnnnnc gtgacagcgc cgccgtcttc gcctggaagg gtgagaccct ccaggagtac 180  
tggtggtgca ccgagcgcgc cctcgactgg ggccccggcg gcggccccga cctncatcgt 240  
cgacgacgga ggcgacgcca ccctcctcat ccacganggc gtcaaggcgg aggagctcta 300  
tgagaagacc ggggaattcc ccgaccctaa ttccantgac aacggcg 347

<210> 3018

<211> 332

<212> nucleic acid

<213> Glycine max

<400> 3018

gtcgcangca cgcntacgtn agctcggaat tcggctcgag caggttctga ccttgaggga 60  
tgttgtttct gaggctgata tctttgtcac caccaccggt aacaaggaca tcatcatggt 120  
tgaccacatg aggaaaatga agaacaatgc cattgtttgc aacattggtc actttgacaa 180  
tgagatcgac atgcttgggc tggagaacta ccccggcgtg aagcgcatca ccatcaagcc 240  
ccaaactgac agatgggtct tccctgagac caacaccggt atcattgtct tggctgaggg 300  
tcgattgatg aacttgggat gcgccactgg ac 332

<210> 3019

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 3019

ngtn gangca cgcgtacgta agctcggaat tcggctcgag gtcaaggacc tntcccaggc 60  
cgacttcggc cgcctcgaga tcgagctggc cgaggttgag atgcccggcc tcatggcctg 120  
tcggacngag ttcgggccct ccagccctt caagggggcc cgcatacccg gctccctcca 180  
catgaccatc cagngcgccg ttctcattga gaccctcacc gcccttggcg ccgaggtccg 240  
ctggtgctcc tgcaacatct tctccacca ggaccacgcc gcgcgcgcta ttgccgcga 300

cagtgcc

307

<210> 3020  
<211> 298  
<212> nucleic acid  
<213> Glycine max

<400> 3020

cgctgcntac gtacgtaagc tcggaattcg gctcgagccg agttcggccc ctcccagccc 60  
ttcaaggggg cccgcatcac cggctccctc cacatgacca tccagaccgc cgttctcatt 120  
gagaccctca ccgcccttgg cgccgaggtc cgctgggtgt cctgcaacat cttctccacc 180  
caggaccacg ccgccgccgc tattgcccgc gacagtgcgc cegtcttcgc ctggaagggt 240  
gagaccctcc aggagtactg gtggtgcacc gagegcgccc tcgactgggg ccccggtg 298

<210> 3021  
<211> 339  
<212> nucleic acid  
<213> Glycine max

<400> 3021

ctttctctag tctgttatt tctcagcgcg taaagcatgg ctttgttggg ggagaaaacc 60  
acgagtggtc gcgantacaa ggtcaaggac tttcccaggc cgacttcggc cgctcgcaga 120  
tcgagctggc cgaggttgan atgcccggcc tcatggcctg tcggaccgag ttcgccccct 180  
cccagccctt caagggggcn cgcatacccg gntccctcca catnaccatc cagaccgcgc 240  
ttctcattga gacctcacc gcccttggcg ccgaggtncg ctggtgctcc tgcaacatct 300  
tctccacca ggaccacgcc gngccgctat tgtcgcgaa 339

<210> 3022  
<211> 275  
<212> nucleic acid  
<213> Glycine max

<400> 3022

caccactggg gttaagaggc tatatcagat gcaggcgaat gggactctac tottccctgc 60  
tattaatgtc aatgactctg ttaccaagag caagtttgac aacttgtaag ggtgccgtca 120  
ctctctccct gatggtctga tgagggtac tgatgtgatg attgctggaa aggtggctgt 180



tggtggctgga tatgggtgatg ttggcaaggg ttgtgctgct gcattgaagc aggctgggtgc 240  
tcgtgtcatc gtgactgaga ttgaccccat ttgtg 275

<210> 3023  
<211> 320  
<212> nucleic acid  
<213> Glycine max

<400> 3023

cnntncacgc gtacgtnagc tcggaattcg gctcaggag gaaaccacca ctggtgttaa 60  
gaggctatat cagatgcagg cgaatgggac tctactcttc cctgctatta atgtcaatga 120  
ctctgttacc aagancaagt ttgacaactt gtacgggtgc cgtcactctc tccctgatgg 180  
tctgatgagg gctactgatg tgatgattgc tggaaagggtg gctgttgtgg ctggatatgg 240  
tgatgttggc aagggttgtg ctgctgcatt gaagcaggct ggtgctcgtg tcatcgtgac 300  
tgagattgac cccattgtgc 320

<210> 3024  
<211> 306  
<212> nucleic acid  
<213> Glycine max

<400> 3024

atgtcgcggtt cacgcgtacg taagctcgga attcggctcg aggggtggaga aaaccacgag 60  
tggtcgcgag tacaaggcca aggacctttc ccaggccgac ttcggccgcc tcgagatcga 120  
gctggccgag gttgagatgc ccggcctcat ggctgtcgg accgagttcg gcccctccca 180  
gcccttcaag ggggcccga tcaccggctc cctccacatg accatccaga ncgcccgttct 240  
cattgagacc ctcaccgccc ttggcgccga ggtccgctgg tgctcctgca acatcttctc 300  
caccca 306

<210> 3025  
<211> 518  
<212> nucleic acid  
<213> Glycine max

<400> 3025

agtttntntt nngggggggg gggggnaang agtancgctn agctatgacg tcgcatgcac 60

gcgtacgtaa gctcggaatt cggctcgagg agagatctat ctatctatca agatggcggt 120  
 gttggttgag aaaacaagca gtggaagga gtacaaggtg aaggacatga cgcaagccga 180  
 ttctggaaga ttggaatcg agctggcgga ggttgaaatg cccggcctca tgtcctccc 240  
 caccgagttc ggccccctctc aacccttcaa gggcgctagg atcaccggct ccctncacat 300  
 gaccatncaa accgncgtcc tcctcgagac cctnaccggc ctngggcgccg aggtcccgt 360  
 ggtgctnctg caacatcttc ttcanccaag accatgccgg cgcgcatcgc cgggacagcg 420  
 ccttcgtctt cgcttgaaa ggtgagacc ttcaggaatc tgggtggtgca ccgagcgcg 480  
 ccttgactgg ggccccngcg gcggcccgat ctattgt 518

<210> 3026  
 <211> 338  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3026

ctccaaggtc agaacctgaa ggccttccat gagagcctga tatctttgtc accaccaccg 60  
 gtaacaagga catcatcatg gttgaccaca tgaggaaaat gaagaacaat gccattgttt 120  
 gcaacattgg tcactttgac aatgagatcg acatgcttgg gctggagaac taccgccggc 180  
 tgaagcgcat caccatcaag ccccaaactg acagatgggt ctncctgag accaacaccg 240  
 gtatcattgt cttggctgag ggtcgattga tgacttggga tgcgccatgg acaccccagt 300  
 tttgtgatgt cctgctcctt caccaacang tcattgtc 338

<210> 3027  
 <211> 286  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3027

gtaccaatgt cggcatcatt gtcttggccg agggctggtt gatgaacttg ggatgcgcca 60  
 caggacaccc tagttttgtg atgtcctgct ccttcaccaa ccaggtcatt gctcagcttg 120  
 agttgtggaa ggagaagagt accggcaagt acgagaagaa agtttacgtt ttgcccgaagc 180  
 acctgatga caaggtggct gcacttcacc ttggcaaact tggagctaag ctcaccaagc 240  
 ttagcccggc ccaggctgat tacatcagtg tgctgttga gggtec 286

<400> 3028

<400> 3029

<400> 3030

1066

tccacatgac catccagacc gccgtttctca ttgagaccct caccgccctt ggcgccgagg 240  
 tccgctggtg ctcttgcaac atcttctcca ccaggacca 280

<210> 3031  
 <211> 324  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3031

ngangcacgc gtacgtaagt cggaattcgg ctcgagtgtt atttctcagc gcgtaaagca 60  
 tgggctttgt tgggtggagaa aaccacgagt ggtcgcgagt acaagggtcaa ggacctttcc 120  
 caggccgact tcggccgcct cgagatcgag ctggccgagg ttgagatgcc cggcctcatg 180  
 gcctgtcgga ccgagttcgg cncctcccag cccttcaagg gggcccgcac caccggctcc 240  
 ctccacatga ccattccagac cgccgttctc attgagacc tcaccgccct tggcgccgag 300  
 ntccgctggt gctctgcaa catc 324

<210> 3032  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3032

acgtcgang caccgctaag tnagctcgga attcggctcg aggcacctcc cagccottca 60  
 agggggcccg catcaccggc tccctccaca tgaccatcca gancgcgct tctcattgag 120  
 accctcaccg cccttggcgc cgaggtccgc tgggtgctct gcaacatctt ctccaccag 180  
 gaccacgccc ccgcgctat tgcccgcgac agtgccgccc tcttcgcctg gaagggtgag 240  
 accctccagg agtactggtg gtgcaccgag cgcgccctcg actggggccc cgggtggtgga 300  
 ccc 303

<210> 3033  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3033

gtnnatncac gcgtacgtaa gtcggaatt cggctcgagg ctggtgctcc tgcaacatct 60

tctccaccca ggaccacgcc gccgccgcta ttgcccgcga cagtgccgcc gtcttcgcct 120  
 ggaaggggtga gaccctccag gactactggt ggtgcaccga gcgcgccctc gactggggcc 180  
 ccggtggtgg acccgacctc atcgtcgacg acggtggtga cgctaccctt ctcatccacg 240  
 aaggcgtcaa ggccgaggag ctctatgaga agaccggcga actccccgac cccaactcca 300  
 ccgacaac 308

<210> 3034  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3034

3034-3035

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 tcatgggtga ccacatgagg aaaatgaaga acaatgccat tgtttgcaac attggtcact 120  
 ttgacaatga gatcgacatg cttggggtgg agaactaccc cggcgtgaag cgcatcacca 180  
 tcaagcccca aactgacaga tgggtcttcc ctgagaccaa caccggatatc attgtcttgg 240  
 ctgagggtcg attgatgaac ttgggatgcg ccaactggaca cccagtttt gtga 294

<210> 3035  
 <211> 332  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3035

ntacagtgcg angcacgcgt acgttagctc ggaattcggc tcgagctcta tcagatgcag 60  
 gcgaatggga ctcttctctt cctgctant aatgtcaatg actctgtcac caangancaag 120  
 ttgacaact tgtatgggtg ccgtcactct ctccctgatg gtctcatgag ggctaccgat 180  
 gttatgattg ctggaaaggt ggctgttggt gctggatatg gtgatgttgg caagggttgt 240  
 gctgctgcaa tgaagcaggc tgggtctcgt gtcacgtga ccgagattga tcccatctgt 300  
 gcccttcagg ctctcatgga agccttcagg tt 332

<210> 3036  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max

<400> 3036

tcncatcgca tgcacgcgta cgtaagctcg gaattcggct cgagcgagta caaggtcaag 60  
gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgcnc 120  
ggcctcatgg nctgtcggac ngagttcggc nctcccagc cttcaaggg ggcccgcatc 180  
accggctccc tccacatgac catccaganc gccgttctca ttgagaccct caccgccctt 240  
ggcgccgagg tccgctggtg ctctgcaac atottctcca ccagga 287

<210> 3037

<211> 326

<212> nucleic acid

<213> Glycine max

<400> 3037

gcacgcgtac gtaagctcgg aattcggctc gaggttccag atcgtgctta ccatcatcag 60  
agatgggttg aagaccgatc ccaccaggta ccgcaagatg aaggagcgtc tcgttgggggt 120  
ttctgaggaa accaccactg gtgttaagag gctatatcag atgcaggcga atgggactct 180  
actcttctg ctattaatgt caatgactct gttaccaaga gcaagtttga caactgttac 240  
gggtgccgtc actctctccc tgatggctctg atgagggcta ctgatgtgat gattgctgga 300  
aagggtggctg ttgtggctgg atatgg 326

<210> 3038

<211> 306

<212> nucleic acid

<213> Glycine max

<400> 3038

cgcangcacg cgtacgtaag ctcggaattc ggctcgaggg aaaatgaaga acaatgccat 60  
tgtttgcaac attggtcact ttgacaatga gatcgacatg cttgggctgg agaactacco 120  
cggcgtgaag cgcatacca tcaagcccca aactgacaga tgggtcttcc ctgagaccaa 180  
caccggtatc attgtcttgg ctgagggctg attgatgaac ttgggatgag ccaactggaca 240  
ccccagtttg tgatgtcctg ctcttcacc aaccaggtea ttgtcagct tgagttgtgg 300  
aaggag 306

<210> 3039

<211> 259  
 <212> nucleic acid  
 <213> Glycine max

<400> 3039

ctcattgaga ccctcaccgc ccttggecgc gaggtccgct ggtgctcctg caacatcttc 60  
 tccacccagg accacgccnc cgccgctatt gcccgcgaca gtgccgccgt ctlogcctgg 120  
 aagggtgaga ccctccagga gtactggtgg tgcaccgagc gcgccctcga ctggggcccc 180  
 ggtggtggac ccgacctcat cgtcgacgac ggtggtgacg ctacccttct catccacgaa 240  
 gcgtcaaggc cgaggagct 259

<210> 3040  
 <211> 306  
 <212> nucleic acid  
 <213> Glycine max

<400> 3040

nncgcatgca cgcntacgta aagctcggaa ttccggtcga ggtaaagcat ggctttgttg 60  
 gtggagaaaa ccacgagtgg tcgogagtac aaggtaagg acctttccca ggccgacttc 120  
 ggccgcctcg agatcgagct ggccgaggtt gagatgcccg gcctcatggc ctgtcggacc 180  
 gagttcggcc cctcccagcc cttcaagggg gcccgcatca ccggctccct ccacatgacc 240  
 atccagancg ccgtttctcat tgagaccctc accgcccttg gcgccgaggt ccgtggtgt 300  
 cctgca 306

<210> 3041  
 <211> 312  
 <212> nucleic acid  
 <213> Glycine max

<400> 3041

ttcangcncg cntacgtaag ctcggaattc ggctcgannc ttgggatgcg ccaactggaca 60  
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 gaaggagaag agtaccggca agtacgagaa gaaggtttac gttttgcca agcaccttga 180  
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gtcaatgact ctgtcaccaa gagcaagttt gacaacttgt atgggtgccg tcaactctctc 180  
cctgatggtc tcatgagggc taccgatgtt atgattgctg gaaaggtggc tgttggtggct 240  
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ctctctccct gatggtctca tgagggctac cgatgttatg attgctggaa aggtggctgt 180  
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 ctctcatggn aggccttcag gntctgacct tggaggatgt tgtttctgan gctgatctct 180  
 ttgtcaccac caccggtaac aaggacatca tcatgggtga ccacatgagg aaaatgaaga 240  
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 ggagaagagt accggcaagt acgagaagaa ggtttacgtt ttgccaagc accttgatga 180  
 gaagggtggct gcacttcacc tgggcaaact tggngctaag ctgaccacgc ttagcaagtc 240  
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 ctaagtgatt gaga 314

<210> 3047  
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cgacttcggc cgcctcgaga tcgagctngc cgaggttgag atgcccggcc tcatggcctg 180  
tcggaccgag ttcgggccct cccagccctt caaggggggc cgcatacccg gctcctccac 240  
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<210> 3048

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<212> nucleic acid

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tgagatcgac atgcttgggc tggagaacta ccccggcgtg angcgcatca ccatcaagcc 180  
ccaaactgac agatgggtct tccctgagac caacaccggt atcattgtct tggtgaggg 240  
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<212> nucleic acid

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cctcgactga ngccccggtg gtggaccgca cctcatcgtc gacgaagggtg gtgacgctac 180  
ccttctcatc cacgaaggcg tcaaggccga ggagctctat gagaagaccg gcgaactccc 240  
cgaccnnaac tccaccgaca acgccgagtt tcagatcggtg cttaccatca tcagagatgg 300  
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 tcaaggacct ttcccaggcc gacttcggcc gcctcgagat cgagctggcc gaggttgaga 180  
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 cccttggcgc cgaggtcgc 319

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tgggactctt ctcttcctg ctattaatgt caatgactct gtcaccaaga gcaagtttga 180

caacttgat gggtgccgtc actctctccc tgatgggtctc atgagggtta ccgatgttat 240

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gtcactttga caatgagatc gacatgctgg ggctggagaa ctaccccggc gtgaagcgca 180

tcaccatcaa gccccaaacc gacagatggg tcttccccga gaccaatgtc ggcatcattg 240

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gagtgggtcgc gagtacaagg tcaaggacct ttcccaggcc gacttcggcc gcctcgagat 180  
 cgagctggcc gaggttgaga tgcccgccct catggnctgt nggaccgagt tcggccccctc 240  
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 acctttccca ggctgacttc ggccgcctcg agatcgagct ggccgaggtc gagatgcccg 180  
 gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg gcccgcatca 240  
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 gcgcnnaggt ccgct 315

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cgaggttgag atgcccggcc tcattggcctg tcggaccgag ttgggcccct cccanccctt 240  
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<212> nucleic acid

<213> Glycine max

<400> 3059

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cggcgtgang cgcataccca tcaagcccca aaccgacaga tgggtcttcc ccgagaccaa 180  
tgtcggcatc attgtcttgg ccgagggctg tttgatgaac ttgggatgcg ccacaggaca 240  
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aggtcaagga ctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180  
agatgcccg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240  
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 gtgctcgtgt catcgtgact gagattgacc ccatttgtgc ccttcaggct ctcatggaag 240  
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 tgggtctcatg agggctaccg atgttatgat tgctggaaaag gtggctgttg tggotggata 240  
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 cccggcgtga agcgcacac catcaagccc caaactgaca gatgggtctt ccctgagacc 180  
 aacaccggta tcattgtctt ggctgagggt cgattgatga acttgggatg cgccactgga 240  
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 g 301

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 tnccacggcc gacttcggcc gcctcgagat cgagctggcc gaggttgaga tgcccggcct 180



catggcctgt cggccgagtt cggccccctcc cagcccttca aggggggccg catcaccggc 240  
 tccctccaca tgaccatcca gaccgcccgtt ctcatgaga ccctcaccgc ccttggcgcc 300  
 gaggtccgct ggtgctctgc aacat 325

<210> 3067  
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 <212> nucleic acid  
 <213> Glycine max  
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cgcngcacgc gtacgtaagc tcggaattcg gctcgagacc gatcccacca ggtaccgcaa 60  
 gatgaaggag cgtctcggtt gggtttctga ggaaaccacc actggtgtta agaggctata 120  
 tcagatgcag gcgaatggga ctctactctt ccctgctatt aatgtcaatg actctgttac 180  
 caagagcaag ttgacaact tgtaogggtg ccgtcactct ctccctgatg gtctgatgag 240  
 ggctactgat gtgatgattg ctggaaangt ggctgttgtg gctggatatg gtgaan 296

<210> 3068  
 <211> 304  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3068

cgtagtaacg tangaanten gntcggtctn aantnntttc tcagcgcgta aagcatggct 60  
 ttgttggtgg anataaccac gagtggtcgc gagtacaagg tcaaggacct ttcccangcc 120  
 gacttcggcc gctcgagat cgagctggcc gangttgana tgcccggcct catggcctgt 180  
 cggaccgagt tcggcccctc ccagcccttc aagggggccc gcatcaccgg ctccctccac 240  
 atgaccatcc agaccgcccgt tntcattgag accctcaccg cccttngngc cgaggnccgc 300  
 tggt 304

<210> 3069  
 <211> 314  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3069

cgcangcacg cgtacgtaag ctcggaattc ggctcgaggc cgtcttccnc tggaagggtg 60

agaccctcca ggagtactgg tgggtgcaccg agcgcgccct cgactggggc cccggtggtg 120  
gacccgacct catcgtcgaa cgacggtggt gacgctaccc ttctcatcca cgaaggcgtc 180  
aaggccgagg agctctatga gaagaccggc gaactccccg accccaactc caccgacaac 240  
gccgagtttc agatcgtgct taccatcadc agagatgggt tgaagaccga tcccaccagg 300  
taccgcaaga tgaa 314

<210> 3070  
<211> 299  
<212> nucleic acid  
<213> Glycine max

<400> 3070

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attgtttgca acattggtca ctttgacaat gagatcgaca tgctggggct ggagaactac 120  
cccgcggtga ngcgcatgca ccatcaagcc ccaaaccgac agatgggtct tccccgagac 180  
caatgtcggc atcattgtct tggccgaggg tcgtttgatg aacttgggat gcgccacagg 240  
acaccctagt tttgtgatgt cctgctcctt caccaaccag gtcattgctc agcttgagt 299

<210> 3071  
<211> 302  
<212> nucleic acid  
<213> Glycine max

<400> 3071

aacgcangca cgcgtagcta agctcngaatt tcggctcgag gtttganaac ttgtacgggn 60  
gccgtcactc tctccctgat ggnetgatga gggctactgt ngtgatgatt gctggaaagg 120  
tggctgttgt ggctggatat ggtgatgttg gcaanggttg tgctgctgca ttgaagcagg 180  
ctggtgctcg tgtcatcgtg actgagattg accccatttg tgcccttcag gctctcatgg 240  
aaggccttca ggttctacct tggaggatgt tgtttctgag gctgatatat ttgtcaccac 300  
ca 302

<210> 3072  
<211> 289  
<212> nucleic acid  
<213> Glycine max

<400> 3072

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gggctctccc cgagaccaat 60  
gtcggcatca ttgtcttggc cgagggctcg ttgatgaact tgggatgcnc cacaggacac 120  
cctagttttg tgatgtcctg ccccttcacc aaccagggtca ttgctcagct tgagttgtgg 180  
aaggagaaga gtaccggcaa gtacgagaag aaagtttacg ttttgcccaa gcaccttgat 240  
gagaagggtg ctgcncttca ccttggcaaa cttgnggcta agctcacca 289

<210> 3073

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 3073

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gcegtcactc tctccctgat ggtctgatga gggctactga tgtgatgatt gctggaaagg 120  
tggtgtttgt ggctggatat ggtgatgttg gcaagggttg tgctgcacca ttgaagcngg 180  
ctggtgctcg tgtcatcgtg actgagattg accccatttg tgcccttcag gctctcttgg 240  
aaggccttca ggttctgacc ttggaggatg ttgtttctga ggctga 286

<210> 3074

<211> 285

<212> nucleic acid

<213> Glycine max

<400> 3074

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gttgggggtt ctgaggaaac caccactgga gttaagaggc tctatcagat gcaggcgaat 120  
gggactcttc tcttccctgc tattaatgct aatgactctg tcaccaagag caagtttgac 180  
aacttgatat ggtgccgtca ctctctccct gatgggtctc tgagggtac cgatgttatg 240  
attgctggaa aggtggctgt tgtggctgga tatgggtgatg ttggc 285

<210> 3075

<211> 300

<212> nucleic acid

<213> Glycine max

<400> 3075

gtcncngcac gcgtacgtna gctcggaatt cggctcgagg ncacgggtaa caaggacatc 60  
atcatgggtg accacatgaa gaaaatgaag aacaatgcc a ttgtttgcaa cattggtcac 120  
tttgacaatg agatcgacat gctggggctg gagaactacc ccggcgtgan gcgcattcac 180  
catcaagccc caaaccgaca gatgggtctt ccccgagacc aatgtcggca tcattgtctg 240  
ggccgagggg cgtttgatga antgggatgc gccacaggac accctagttt tgtgatgtcc 300

<210> 3076

<211> 264

<212> nucleic acid

<213> Glycine max

<400> 3076

ccgaggtcog ctggtgctcc tgcaacatct tctccacca ggaccacgcc gccgccgcta 60  
ttgccgcna cagtgccgcc gtcttcnct ggaagggtga gacctccag gactactggt 120  
ggtgcaccga gcgcgccctg cgactggggc cccggtggtg gaccgcacct catcgtcgan 180  
nacggtggtg acgctaccct tctcatccag gaaggcgtca aggcgcagga gctctatgag 240  
aagaccggcg aactccccga ncct 264

<210> 3077

<211> 310

<212> nucleic acid

<213> Glycine max

<400> 3077

ngcangcacg cgtacgtaan ctcggaattc ggctcgagta gtccgtgttat ttctcagcgc 60  
gtaaagcatg gctttgttgg tggagaaaac cacgagtggt gcgcagtaca aggtcaanga 120  
cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg agatgcccg 180  
cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg ccgcacac 240  
cggctccctc cacatgacca tenantcaaa ngttctcatt gagaccctca ccgcccttgg 300  
cgccgaggtc 310

<210> 3078

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 3078

ncnatgcacg cgtacgtaag ctcggaattc ggctcgagct caccactccc tccactctct 60  
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 gagtggtcgc gagtacaagg tcaaggacct ttcccaggcc gacttcggcc gcctcgagat 180  
 cgagctggcc gaggttgaga tgcccggcct catggcctgt cggaccgagt tcggcccctc 240  
 ccagcccttc aagggggccc gcatcaccgg ctccctccac atgaccatcc agancgccgt 300  
 totcattgag accctcaccg ccctt 325

<210> 3079

<211> 307

<212> nucleic acid

<213> Glycine max

<400> 3079

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 tgctattaat gtcaatgact ctgtcaccan gagcaagttt gacaacttgt atgggngccg 180  
 tcactctctc cctgatggtc tcatganngc taccgatntt atgattgctg gaaaggtggc 240  
 tgttggtggn ggatatggtg atgttggnan gggttgtgct gctncaatnn agcaggctgg 300  
 tgetenc 307

<210> 3080

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 3080

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 gcgtaaagca tggctttgtt ggtggagaaa accacgagtg gtcgcgagta caaggtcaag 120  
 gacctttccc aggcgcactt cggccgcctc gagatcgagc tggccgaggt tgagatgcc 180  
 ggctcatgg cctgtcggac cgagttcggc cctcccagc ccttcaaggg ggcccgcac 240  
 accggctccc tccacatgac catccaganc gccgtttctc ttgagacct caccgccctt 300

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<210> 3081  
<211> 293  
<212> nucleic acid  
<213> Glycine max

<400> 3081

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagg ttattttctca gcgcgtaaag 60  
catggctttg ttggtggaga aaaccacgag tggtcgogag tacaaggta aggacctttc 120  
ccaggccgac ttcgcccgcc tcgagatcga gctggccgag gttgagatgc ccggcctcat 180  
ggcctgtcgg accgagttcg gcccctccca gcccttcaag ggggcccgc taccggctc 240  
cctccacatg accatccaga ccgcggttct cattgagacc ctcaccgccc ttg 293

<210> 3082  
<211> 309  
<212> nucleic acid  
<213> Glycine max

<400> 3082

agtcgcatgt ntagtacgta agctcggaat tcggctcgag ctctagtcct gttattttctc 60  
agcgcgtaaa gcatggcttt gttggtggag aaaaccacga gtggtcgca gtacaaggta 120  
aaggaccttt ccagggcga ctctggccgc ctcgagatcg agctggccga ggtcgagatg 180  
cccggcctca tggcctgttc ggaccgagtt cggcccctcc cagcccttca agggggcccg 240  
catcaccggc tccctccaca tgaccatcca gaccgcggtt ctcattgaga ccctcaccgc 300  
ccttggcgc 309

<210> 3083  
<211> 295  
<212> nucleic acid  
<213> Glycine max

<400> 3083

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gcgtaaagca tggctttggt ggtggagaaa accacgagtg gtcgcgagta caaggtaag 120  
gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgccc 180

ggcctcatgg cctgtcggac cgagttcggc ccctcccagc cttcaaggg ggcccgcac 240  
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<210> 3084  
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ggacctttcc caggccgact tcggccgcct cgagatcgag ctggccgagg ttgagatgcc 180  
cggcctcatg gcctgtcggg cagagttcgg ccctcccag cccttcaagg gggcccgcac 240  
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gcgtaaagca tggctttgtt ggtggagaaa accacgagtg gtcgcgagta caaggtcaag 120  
gacctttccc aggcgcgactt cggccgcctc gagatcgagc tggccgaggt tgagatgcc 180  
ggcctcatgg cctgtcggac cgagttcggc ccctcccagc cttcaaggg ggcccgcac 240  
accgggtccc tccacatgac catccagacc gccgtttctca ttgagaccct cac 293

<210> 3086  
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cgagtggctcg cgagtacaag gtcaaggacc tttcccaggc cgacttcggc cgcctcgaga 180  
 tcgagctggc cgagggttgag atgcccggcc tcatggcctg tcggaccgag ttcgggccct 240  
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 ttctcattga gacctcacc gc 322

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 aaagtattacg ttttgcccaa gcaccttgat gagaagggtg ctgcacttca ccttggcaaa 180  
 cttggagcta agctcaccaa gcttagcccg gccaggctg attacatcag tgtgcctgtt 240  
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<210> 3088  
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 <212> nucleic acid  
 <213> Glycine max  
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 cgagctggcc gaggttgaga tgcccggcct catggcctgt cggaccgagt tcggccctc 240  
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<210> 3089  
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 <212> nucleic acid  
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<210> 3090  
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<400> 3090

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 cctcaccct tggcgccg 318

<210> 3091  
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 <212> nucleic acid  
 <213> Glycine max

<400> 3091

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 aggttgagat gcccggcctc atggcctgtc ggaccgagtt cggcccctcc cagcccttca 180  
 agggggcccg catcaccggc tccctccaca tgaccatcca gaccgcccgtt ctcattgaga 240  
 ccctcaccgc cttggcggn gacnncgggn nctnaaaaa 279

<210> 3092  
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 <212> nucleic acid

<213> Glycine max

<400> 3092

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acctttccca ggccgaacttc ngccgcctcg agatcgagct ggccgagggt gagatgcccg 180  
gctcatggc ctgtcggacc gagttoggcc cctcccagcc cttcaagggg gcccgcatca 240  
cgggtccct ccanatgacc atccagaccg ccgttctcat tgagacnctc accgcccttg 300  
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<210> 3093

<211> 242

<212> nucleic acid

<213> Glycine max

<400> 3093

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gcatcaccat caagcccaaa accgacagat gggctctccc gagaccaatg tcggcatcat 180  
tgtctggcgg agggctgttt gatgaacttg ggatgcgcca caggacaccc tagttttgtg 240  
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<210> 3094

<211> 303

<212> nucleic acid

<213> Glycine max

<400> 3094

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ctcagcgct aaagcatggc tttgttggtg gagaaaacca cgagtggctc cgagtacaag 120  
gtcaaggacc tttcccaggc cgacttcggc cgctcagaga tcgagctggc cgaggctcag 180  
atgcccgccc tcatggcctg tcggaccgag ttccggccct cccagccctt caagggggcc 240  
cgcatcaccg gtcctccac atgaccatcc agaccgccgt tctcattgag accctcaccg 300  
ccc 303

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agcgcgtaan gcntggcttt gttggtggag aaaaccacga gtggtcgcga gtncaaggtc 120  
naggnccttt cccaggccgn cttcggccgc ctcgagatcg agctggccga ggttgagnnq 180

cccggcctca tggcctgtcg gaccgagttc ggccccctccc ancccttcaa gggggcccgc 240  
 atcaccggct cctccacat gacntccag accgccgttc tcattgagac cctcancgcc 300  
 cttggcgcc 309

<210> 3098  
 <211> 272  
 <212> nucleic acid  
 <213> Glycine max  
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 cgttgggggtt ttgaggaaa ccaccactgg agttaagagg ctctatcaga tgcaggcgaa 120  
 tgggactctt ctcttccctg ctattaatgt caatgactct gtcaccaaga gcaagtttga 180  
 caacttgat gggtgccgtc actctctccc tgatgggtctc atgagggcta ccgatgttat 240  
 gattgctgga aagggtggctg ttgtggctgg at 272

<210> 3099  
 <211> 339  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3099

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 cgaggttgag atgcccgcc tcatggcctg tcggncggag ttcggcncct cccagccctt 180  
 caagggggcn cgcataccg gntccctcca catgaccatc cagancgccg ttctcatttg 240  
 agatcctnat cgcccttggn gccgnaggtc cgctgggtgt cctgnaacat cgtctccatc 300  
 caggaccacg ccncngccgc tattgcccg anagtgcg 339

<210> 3100  
 <211> 262  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3100

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cgcgagtaca aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg 120  
 gccgaggttg agatgcccg cctcatggcc tgtcggaccg agttcggccc ctcccagccc 180  
 ttcaaggggg ccgcgcatcac cggtccctc cacatgacca tccagctccg ccgtttctcat 240  
 tgagaccctc accgcccttg gc 262

<210> 3101  
 <211> 276  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3101

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 gcaggttgtt gctcgtgtca tcgtgactga gattgacccc atttnggccc ttnaggttct 120  
 catggaaggc cttcaggttc tgaccttgga ggatgttggt tctgaggctg atatctttgt 180  
 caccaccaag ggtaacaagg acatcatcat ggttgaccac atgaagaaaa tgangancan 240  
 tgccattgtt tgcaacattg gtcactttga caatga 276

<210> 3102  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3102

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 ttctcagcgc gtaaagcatg gctttnttgg tggagaaaac caccagtggt cgcgagtaca 120  
 aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180  
 agatgcccg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240  
 ccgcgcatcac cggtccctc cacatgacca tccagaccgc cgttctcatt gagacc 296

<210> 3103  
 <211> 294  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3103

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tttcacatga ccatccagac cgtgtgtoctc atcgagaccc tcaccgctct cggcgccgag 120  
gttcgctggt gtcctgcaa catcttctcc actcaggacc acgcccgcgc cgccatcgcc 180  
cgtgacagcg ccgcgctctt cgootggaag ggtgagaccc tccaggagta ctggtngtgc 240  
accgagcncg cctcgactg gggccccggc ggcgcccccg acctcatcgt cgac 294

<210> 3104  
<211> 291  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3104

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gcgtaaagca tggctttgtt ggtggagaaa accacgagtg gtcgcgagta caaggtcaag 120  
gacctttccc aggcgactt cggccnctc gagatcgagc tggccgaggt tgagatgccc 180  
ggcctcatgg cctgtcggac cgagttcggc cctcccagc cttcaaggg ggcccgcac 240  
accggctccc tccacatgac catccagacc gccgttctca ttgagaccct c 291

<210> 3105  
<211> 311  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3105

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ggtggtgacg ctacccttct catccacgaa ggctcaagg ccgaggagct ctatgagaag 120  
accggcgaac tccccgaccc caactccacc gacaacgccg agtttcagat cgtgcttacc 180  
atcatcagag atggggtgaa gaccgatccc accagggtacc gcaagatgaa ggagcgtctc 240  
gttgggggttt ctgaggaaac caccactgga gttaagaggc tctatcagat gcaggcgatt 300  
gggccttttn t 311

<210> 3106  
<211> 301  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3106

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 ccgacttcgg ccgcctcgag atcgagctgg ccgaggttga gatgcccggc ctcatggcct 180  
 gttcggaccg agttcggccc ctcccagccc ttcaaggggg cccgcacac cggctccctc 240  
 cacatgacca tccagaccgc cgtttctcatt gagaccctca ccgccttgnc gccgaggtcc 300  
 g 301

<210> 3107  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max

<400> 3107

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 agaggctcta tcagatgcag gcgaatggga ctcttctctt ccctgctatt aatgtcaatg 120  
 actctgtcac caagagcaag ttgacaact tgtatgggtg ccgtcatctc tccctgatgg 180  
 tctcatgagg gctaccgatg ttatgattgc tggaaagggtg gctgttgtgg ctggatatgg 240  
 tgatgttggc aagggttgtg ctgctgcaat gangcagggtg gtccccgttc a 291

<210> 3108  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max

<400> 3108

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 cgagcgcgcc ctcgactggg gccccgggtg tggacccgac ctcatcgctg acgacgggtg 120  
 tgacgctacc cttctcatcc acgaaggcgt caaggccgag gagctctatg agaaaccggc 180  
 gaactccccg accccaactc caccgacaac gccgagattc agatcgtgct taccatcatc 240  
 agagatgggt tgaagaccga tcccaccagg taccgcaaga tgaaggagcg tctcgttg 298

<210> 3109  
 <211> 341  
 <212> nucleic acid  
 <213> Glycine max

<400> 3109

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acaatgccat tgtttgcaac attgggtcact ttgacaatga gatcgacatg ctgggggctgg 120  
agaactaccc cggcgtgaag cgcataacca tcaagcccca aacngacaga tgggtcttcc 180  
ccgagaccaa tgtcggcatc attgtcttgg ccgagggctcg tttgatgaac ttgggatgcg 240  
ccacaggaca ccctagtttt gtgatgtctg tnccttcacc aaccagggtca tgctcagttg 300  
agttgtggaa angagaagag taccggcaag tacgagaagn a 341

<210> 3110

<211> 279

<212> nucleic acid

<213> Glycine max

<400> 3110

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ctccaccag gaccangcgc ccgcgcgtat tgcccgcgac agtgcngccg tcttcgcctg 120  
gaaggggtgan accctccagg agtactggtg gtgcaccgag cgcgccctcg actggggccc 180  
cggtggtgga ccgacctca tcgtcgacga cggtggtgac gctacccttc tcatccacga 240  
aggcgtcaag gccgaggagc tctatnagaa gaccggcga 279

<210> 3111

<211> 271

<212> nucleic acid

<213> Glycine max

<400> 3111

ccaccangta ccgcaagatg aaggagcgtc tcgttgggggt ttctgaggaa accaccactg 60  
gtgttaanan gctatatcag atncaggcna atgggantct actcttccct gctattaatg 120  
tcaatgactc tgttaccaag agcaagtttg acaacttgta cgggtgccgt cactctctcc 180  
ctgatggtct gatgagggct actgatgtga tgattgctgg aaaggtggct gttgnggcc 240  
ggatanggtg atnttggcaa gggttngcn c 271

<210> 3112

<211> 293

<212> nucleic acid



3112  
3113  
3114  
3115

<213> Glycine max

<400> 3112

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tcagcgcgta aagcatggct ttgttggtgg agaaaaccac gantggtcgc gagtacaagg 120  
tcaaggacct ttcccaggcc gacttcggcn gcctcgagat cgagctggcc gaggttgaga 180  
tgcccggcct catggcctgt cggaccgagt tcggcccctc ccagcccttc aagggggccc 240  
gcatcaccgg ctcccctcac atgaccatcc agaccgccgt tctcattgag acc 293

<210> 3113

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 3113

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atctctcagc gcgtaaagca tggctttgtt ggtggagaaa accacgagtn gtcgcgagta 120  
caagggtcaag gacctttccc aggcgcactt cggccgcctc gagatcgagc tggccgaggt 180  
tgagatgcc ggctcatgg cctgtcggac cgagttcggc cctcccagc ccttcanggg 240  
ggcccgcatc accggtctcc tccacatgac catccagacc gccgtttctca ttgagaccct 300  
n 301

<210> 3114

<211> 283

<212> nucleic acid

<213> Glycine max

<400> 3114

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cccttcaagg gggcccgcat caccggcncc cttcacatga ccatcnagac cgtgtctctc 180  
atcgagacct tcaccgtct cggcgccgag gttcgtggt gtcctgcaa catcttctcc 240  
actcaggacc acgccgccgc cgccatcgcc cgtgacagcg ccg 283

<210> 3115

<211> 313  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3115  
  
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 aggtcaagga cctttcccag gcgacttcg gccgcctcga gatcgagctg gccgaggttg 180  
 anatgcccgg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240  
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 ngcccttggg cga 313

<210> 3116  
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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3116  
  
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 acaaggncna ggacctttcc caggccgact tcggccgcct cgagatcgag ctggccgagg 180  
 ttgagatgcc cggcctcatg gcctgtcggg ccgagttcgg cccctcccag cccttcaagg 240  
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 caccg 305

<210> 3117  
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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3117  
  
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 gcatggcttt gttggtggag aaaaccacga gtggctcga gtacaaggtc aaggaccttt 120  
 cccaggccga cttcggccgc ctccagatcg agctggccga ggttgagatg cccggcctca 180  
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ccctccacat gaccatccag acagccgttc tcattgaga

279

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<213> Glycine max  
  
<400> 3118

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gcgcgtaaag catggctttg ttggtggaga aaaccacgag tgggtgcgca gtacaaggtc 120  
aaggaccttt cccaggccga cttcggecgc ctcgagatcg agctggccga ggttgagatg 180  
cccggcctca tggcctgtcg gaccgagttc ggccccctcc agcccttcaa gggggcccgc 240  
atcacgggct ccctccacat gaccatccag accgccgttc tcattgagac ctcaccgcc 300  
c 301

<210> 3119  
<211> 322  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3119

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tcanggnagg ccttcaggnt ctgaccttng aggatgttng ttctgaggct gatatcngtg 120  
tcaccancca nccgtaacaa ggacatcatc atgggtgacc acatgangan aatgaagaac 180  
aatgccattg tttgcaacat tggctcattg acaatgagat cgacatgctt gggctggaga 240  
actaccccg cgtgaagcgc atcaccatca agccccaac tgacagatgg gtcttccctg 300  
agaccaaacac cggatcatgt ct 322

<210> 3120  
<211> 293  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3120

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tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120



aanagcatgc acgcgtacgt aagctcggaa ttcggctcga gccactctct ttctctagtc 60  
 ctgttatttc tcagcgcgta aagcatggct ttgttggtgg agaaaaccac gagtggtcgc 120  
 gagtacaagg tcaaggacct ttcccaggcc gacttcggcc gcctcgagat cgagctggcc 180  
 gaggttgaga tgcccggcct catggcctgt cggaccgagt tcggcccctc ccagcccttc 240  
 aagggggccc gcatcaccgg ctccctccac atgaccatcc agaccgccgt tctcatt 297

<210> 3124  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3124

gtcgcangca cgcgtacgta agctcggaa ttcggctcga ctctttctct agtcctgtta 60  
 tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120  
 aaggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgaggtt 180  
 gagatgcccg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240  
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<210> 3125  
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 <400> 3125

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 accccggcgt gangcgcac accatcaagc cccaaaccga cagatggtct tccccgagac 180  
 caatgtcggc atcattgtct tggccgaggg tcgtttgatg aacttgggat gcgccacagg 240  
 acaccctagt tttntgatgn cctgctcctt cac 273

<210> 3126  
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 <212> nucleic acid  
 <213> Glycine max  
 <400> 3126

ctnccgcangc ncgcgtacgt aagctcggaa ttcggctcga gctctttctc tagtcctggt 60  
 atttctcagc gcgtaaagca tggctntggt ggtggagaaa accacgagtg gtcgcgngta 120  
 caaggccaag gacctttccc aggcgcactt cggccgcctc gagatcgagc tggccgaggt 180  
 tgagatgcc ggccatcatgg cctgtcggac cgagttcggc ccctcccagc cttcaagggt 240  
 ggcccgcatc accggtccc tccacatgac catccaganc gccgttctc 289

<210> 3127  
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 <212> nucleic acid  
 <213> Glycine max

<400> 3127

3127

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 caaggacctt tcccaggccg acttcggncg cctngagatc ganctggccg aggttganan 180  
 gcacggcctc atggcctgtc ggaccgagtt cngcccctcc cancccttca agggggcccg 240  
 catcaccggc tccctccaca tgaccatcca gaccggcggtt ctcatcgaga ccctcaccgn 300  
 cttggcgccg 310

<210> 3128  
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 <212> nucleic acid  
 <213> Glycine max

<400> 3128

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 cgggtggtgga cccgacctca tcgtcgacga cgggtggtgac gctacccttc tcatccacga 120  
 aggcgtcaag gccgaggagc tctatgagaa gaccggcgaa ctccccgacc ccaactccac 180  
 cgacaacgcc gagtttcaga tcgtgcttac catcatcaga gatgggttga agaccgatcc 240  
 caccaggtac cgcaagatga aggagcgtct cgttgggggtt tctgaggaaa cc 292

<210> 3129  
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 <212> nucleic acid  
 <213> Glycine max

<400> 3129

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tctcagcgcg taaagcatgg ctttggttgg ggagaaaacc acnctgtgtg cgcgagtaca 120  
aggtcaagga cctttcccag gccgacttcg gccgcctcga gatcgagctg gccgaggttg 180  
agatgcccgg cctcatggcc tgtcggaccg agttcggccc ctcccagccc ttcaaggggg 240  
cccgcatac cggctccctc cacatgacca tccagancgc cgttctcatt gagaccctc 299

<210> 3130

<211> 325

<212> nucleic acid

<213> Glycine max

<400> 3130

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ttctcagcgc gtaaagcatg gctttgttgg tggagaaaac cacgagtggc ccgcgagtac 120  
aaggtcaagg acctttccca gccgacttc gccgcctcga agatcgagct gccgaggtt 180  
gagatgcccg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240  
gccgcatac cggctccct ccacatgacc atccagaccg ccgttctcat tgagacctca 300  
ccgccttggc gccgaggtcg cttnn 325

<210> 3131

<211> 273

<212> nucleic acid

<213> Glycine max

<400> 3131

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ttggtggaga aaaccacgag tggtcgcgag tacaaggta aggaccttc ccaggccgac 120  
ttcggccgcc tcgagatcga gctggccgag gttnagatgc ncggcctcat ggctgtcgg 180  
accgagttcg gnnctncca gcccttcaag ggggcncgca tcanoggntc cctccacatg 240  
accatcnagn ccgccgttct cattgagacc etc 273

<210> 3132

<211> 286

<212> nucleic acid

<213> Glycine max

<400> 3132

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aagggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgagggtt 180  
gagangcccg gctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240  
gcccgcata cgggtccct ccacatgacc atccagaccg ccgttc 286

<210> 3133

<211> 288

<212> nucleic acid

<213> Glycine max

<400> 3133

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aagggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgagggtt 180  
gagangcccg gctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240  
gcccgcata cgggtccct ccacatgacc atccagatcg ccgttctc 288

<210> 3134

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 3134

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tttcccaggc cgacttcggc cgctcggaga tcgagctggc cgaggttgag atgcccggcc 180  
tcattggcctg tcggaccgag ttccggcccct ccagcccctt caagggggcc cgcatacccg 240  
gtccctcca natgaccatn cagaccgccc tccctcattg agaccctca 289

<210> 3135

<211> 289

<212> nucleic acid



6466766

<213> Glycine max

<400> 3135

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 cgctaccctt ctcatocacg aaggcgtcaa ggccgaggag ctctatgaga agaccggcga 120  
 actccccgac cccaactcca ccgacaacgc cgagtttcag atcgtgctta ccatcatcag 180  
 agatgggttg aagaccgatc ccaccaggta ccgcaagatg aaggagcgtc tcgttggggt 240  
 ttctgaggaa accaccactg gagttaagag gctctatcag atgcaggcg 289

<210> 3136

<211> 281

<212> nucleic acid

<213> Glycine max

<400> 3136

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 gognaaagca tggctttgtt ggtggagaaa accacgagtg gtcgagagta caaggccaag 120  
 gacctttccc agnccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgccc 180  
 ggccatcatgg cctgtcggac cgagttcggc ccctcccagc ctttcaaggg ggcccgcac 240  
 accggctccc tccacatgac catccatgac accgtttctca t 281

<210> 3137

<211> 301

<212> nucleic acid

<213> Glycine max

<400> 3137

gtcgctgcac gcgtacgtaa gctcggaatt cggctcgagc tagttttgtg atgtcctgct 60  
 cnttcaccan nccaggatcat tgctcagctt gagttgtgga aggagaagag taccggcaag 120  
 tacgagaaga aagtttacgt ttgcccgaag caccttgatg agaaggtggc tgcacttcac 180  
 cttggcaaan ttggagctaa gtcaccaag cttagcccgg ccagggctga ttacatcagt 240  
 gtgcctgttg agggccata caagcctgct cattacaggt actaagtaat tgagattatc 300  
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 <212> nucleic acid  
 <213> Glycine max

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 aaggtaagg acctttccca ggccgacttc ggccgctcg agatcgagct ggccgagggt 180  
 gagatgcccg gctcatggn ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240  
 gnccgcatca ccggtccct ccacatgacc atccagancg ccgttn 286

<210> 3139  
 <211> 270  
 <212> nucleic acid  
 <213> Glycine max

<400> 3139

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 tgctggaaag gtggctgttg tggctggata tggatgatgt ggcaaggggt gtgctgctgc 180  
 attgaagcag gctggtgctc gtgtcatcgt gactgagatt gacccattt gtgccottca 240  
 ggctctcatg gaaggcctca gttctgacct 270

<210> 3140  
 <211> 298  
 <212> nucleic acid  
 <213> Glycine max

<400> 3140

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 ggtggtgcac cgagcgcgcc ctacgactgg ggccccggtg gtggaccgga cctcatcgtc 120  
 gacgacggtg angacgtac cttctcatc cacgaaggcg tcaaggccga ggantctatg 180  
 agacgaccgg cgaactcccc gaccccaact ccaccgacaa cgccgagttt cagatcgtgc 240  
 ttaccatcat cnganatggg ttgaagaccg atcccaccag gtaccgcaag atgaagga 298

<210> 3141

<211> 334  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3141  
  
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 agttaagagg ctctatcaga tgcnaggcga atgggattcc tctcttccng ntaataaatg 120  
 tcaatgactc ntgtcaccan gagcnagttt gacaacttgt atgggtgccg tncactctct 180  
 ccctgatggt ctcatgaggg ctaccgatgt tatgattgct ggaaagggtg ctgttgtngc 240  
 tggatatggt gatgttggca anggttgtgc tgctgcaatg naggaggctg gtgctcgtgt 300  
 catcgtgnac gagattgac ccattctgtgc cctc 334

<210> 3142  
 <211> 266  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3142  
  
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 aaccacgagt ggtcgcgagt acaagggtcaa ggacctttcc caggccgact tcggccgcct 120  
 cgagatcgag ctggccgagg ttgagatgcc cggcctcatg gcctgtcgga ccgagttcgg 180  
 cccctcccag ccttcaagg gggcccgcat caccggtcc ctccacatna cnaanaaatn 240  
 ncnaantctc attgagaccc tcancg 266

<210> 3143  
 <211> 288  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3143  
  
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 caaggacctt tcccaggccg acttcggccg cctcgagatc gagctggccg aggttgagat 180  
 gcccggcctc atggcctgtc ggaccgagtt cggccctcc cagcccttca agggggcccg 240  
 catcaccggc tcctccaca tgaccatcca ganncgccgt tctcattg 288

<210> 3144  
 <211> 308  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3144  
  
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 tctcatccac gaaggcggtca aggccgagga gctctatgag aagaccggcg aactccccga 120  
 ccccaactcc accgacaacg ccgagtttca gatcgtgctt ancatcatca gagatggggtt 180  
 gaagaccgat cccaccaggt ancgcaagat gaaggagcgt ctcggtgggg tttctgagga 240  
 aaccaccatt ggagttaaga ggntctatca gatgcaggcg aatgggatct tctcttcctt 300  
 gctattaa 308

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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3145  
  
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 atttctcagc gcgtaaagca tggctttggt ggtggagaaa accacgagtg gtcgcgagta 120  
 caaggtaag gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt 180  
 tgagatgcc ggctcatgg cctgtcggac cgagttcggc cctcccagc cttcaaggg 240  
 ggcccgcatc accggtccc tccacatgac catccagac 279

<210> 3146  
 <211> 296  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3146  
  
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 tccacgaggg cgtcaaggcc gaggagctct atgagaagac cggggaactc cccgacccta 120  
 actccactga caacgccgag ntccagatcg tgcttaccat catcagagat gggttgaaga 180  
 ccgatcccac caggtaccgc aagatgaagg agcgtctcgt tggggtttct gaggaacca 240

ccactggtgt taagaggcta tatcagatgc aggcgaatgg gactctactc ttccct 296

<210> 3147  
 <211> 287  
 <212> nucleic acid  
 <213> Glycine max

<400> 3147

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gtggagaaaa ccacgagtgg tcgcgagtta caaggtcaag gacctttccc aggccgactt 120

cggccgcctc gagatcgagc tggccgaggt tgagatgccc ggcctcaggc ctgttcggac 180

cgagttcggc cctcccagc cttcaaggg ggccgcgcatc accggtccc tccacatgac 240

catccagacc gccgtttctca tgagaccctc accgcccttg gcgccga 287

3148  
 275  
 nucleic acid  
 Glycine max  
 3148

<210> 3148  
 <211> 275  
 <212> nucleic acid  
 <213> Glycine max

<400> 3148

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tttctcagcg cgtaaagcat ggctttgttg gtggagaaaa ccacgagtgg tcgcgagtac 120

aaggtcaagg acctttccca ggccgacttc ggccgcctcg agatcgagct ggccgaggtt 180

gagatgcccg gctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240

gcccgcacga ccggctccct ccacatgacc atcca 275

<210> 3149  
 <211> 239  
 <212> nucleic acid  
 <213> Glycine max

<400> 3149

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gttgtggatg gagaagagta ccggcaagta cgagaagaag gtttacgttt tgcccaagca 120

ccttgatgag aaggtggctg cacttcacct gggcaaactt ggngctaagc tgaccagct 180

tagcaagtcc caggctgatt acatcagtg gctgttgag ggtccatata agcctgctc 239

<210> 3150  
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 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3150  
  
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 ggtcaaggac ctttcccagg ccgacttcgg ccgcctcgag atcgagctgg ccgagggtga 180  
 gatgcccggc ctcatggcct gtcggaccga gttcggcccc tcccagccct tcaagggggc 240  
 ccgcatacc ggctccctcc acatgaccat 270

<210> 3151  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3151  
  
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 gtggagaaaa ccacgagtgg tcgcgagtac aagggtcaagg acctttccca ggccgacttc 120  
 ggccgcctcg agatcgagct ggccgagggt gagatgcccg gcctcatggc ctgtcggacc 180  
 gagttcggcc cctcccagcc cttcaagggg gcccgcatca ccggtccct ccacatnaca 240  
 nnnnacngaa aaatgctcat tgagaccctt caccgccnnt gggggcgngg 290

<210> 3152  
 <211> 310  
 <212> nucleic acid  
 <213> Glycine max  
  
 <400> 3152  
  
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 cgagtacaag gtcaaggacc tttcccaggc cgacttcggc cgctcgaga tcgagctggc 180  
 cgagggttgag atgcccggcc tcatggcctg tcggancgag ttccggccct cccagccctt 240  
 caagggggcc cgcatacccg gctccctcca natgaccatc cagaccgccg ttctcattga 300

gagctcacgc 310

<210>	3153
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<212>	nucleic acid
<213>	Glycine max

<400> 3153

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cgcgtaaaagc	atggcttttgt	tgggtggagaa	aaccacnagt	ggtacgcgag	tacaagggtca	120
aggacaccttc	ccaggcccgac	ttcggcgcgcc	tcgagatcga	gctggccgag	gttgagatgc	180
ccggcctcat	ggcctgtcgg	accgagttcg	gccctccca	gcccttcaag	ggggcccgca	240
tcaccgggctc	cctccacatg	accatccaga	ccgcgcgt			277

[illegible]

<210>	3154
<211>	298
<212>	nucleic acid
<213>	Glycine max

<400> 3154

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ttctcagcgc	gtaaagcatg	gctttgttgg	tggagaaaac	cacgagtggg	cgcgagtaca	120
aggtcaagga	cctttcccag	gcgacttcg	gccgcctoga	gategagctg	gccgaggttg	180
agatgcccgg	cctcatggcc	tntcggaacc	agttngggcc	cgncagccc	gtnaaggggg	240
cccgcacnc	cggcgctcgc	nacaggatca	nccagaccgc	cgttctcagt	ganacccg	298

<210>	3155
<211>	318
<212>	nucleic acid
<213>	Glycine max

<400>	3155
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cgctccactc	tctttctcta	gtcctgttat	ttctcagcgc	gtaaancatg	gctnanttgg	120
tggagaaaac	cacgagtgg	cgcnagtaca	aggtaagga	cctttcccag	gccgacttcg	180
gcgcctcgag	atcgagctgg	cgcnggttga	gatgcccggc	ctcatggcct	gtncggaccg	240

agttcgcccc ctcccagccc ttcaaggggg cccgcatcac cggtccctc cacatgacca 300  
tccagaccgc cgtttctca 318

<210> 3156  
<211> 318  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3156

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aaaaccacga gtggtcgcca gtacaaggtc aaggaccttt ccagggccga cttcggccgc 120  
ctcgagatcg agctggccga ggttgagatc ccggcctcat ggctgtngg accgagttcg 180  
gcccctcccag cctcaaggg ggcccgcac accggctccc tccacatgac catccagacc 240  
gccgtttctca ttgagacctc acngccttgg gccgagtcgg ttggtgtctt gaaanatttc 300  
tcaaccaagg acaagcng 318

<210> 3157  
<211> 292  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3157

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tctcngcgcg taaagcatgg ctttgttggg ggagaaaacc angagtgggc gcgagtacaa 120  
ggtcaaggac gtttcccagg ccgacttcgg ccgcctcgag atcgagctgg ccgngggtga 180  
gatgcccggc ctcatggcct gtcggaccga gttcggcccc tcccagccct tcaagggggc 240  
ccgcatcacc ggctccctcc acatnacann cgacngcanc gttctcattg an 292

<210> 3158  
<211> 278  
<212> nucleic acid  
<213> Glycine max  
  
<400> 3158

tcgcangcac gcgtacgtna gtcggaatt cggtcagagn tttctctagt cctgttattt 60  
ctcagcgcgt aaagcatggc tttgttgggt gagaaaacca cgagtggtcg cgagtacaag 120



gtcaaggacc tttcccaggc cgacttcggc cgctcgcaga tcgagctggc cgaggttgag 180  
atgcccggcc tcatggcctg tcggaccgag ttcgggccct cccagccctt caagggggcc 240  
cgcatcaccg gctccctcca catggaccat ccagaccg 278

<210> 3159  
<211> 332  
<212> nucleic acid  
<213> Glycine max

<400> 3159

acaccncct accacgccaan cgnaagctcg gaattnggct cgagattcac caaccaggt 60  
cattgctcag ttgagttgtg gaaggagnag agtaccggca agtacgagaa gaaggtttac 120  
gttttgcnca agcaccttga tgagaagggtg gctgcactta acctgggcaa acttgnagct 180  
aagctgaccc agcttagcaa gtnccaggnt gattacatca gtgtgcctgt tgaggggtcca 240  
tacaagcctg ctcantacag gtacnnnctn atnnngatga tcaactgnaa agtgagtgag 300  
ggaaagacaa aaatgggttt tatnaatngg at 332

<210> 3160  
<211> 288  
<212> nucleic acid  
<213> Glycine max

<400> 3160

tcgcangcac gcgtacgtaa gctcggaatt cggctcgagc tctttctcta gtcctgttat 60  
ttctcagcgc gtaaagcatg gctttgttgg tggagaaaac cacgagtggc cgcgagtaca 120  
aggtcaagga cttttccag gccgacttcg gccgcctcga gatcgagtgg ccgaggttga 180  
gatgcccggc ctcatggcct ntcggaccga gttcgggccc tcccagccct tcaagggggc 240  
ccgcatcacc ggctccctcc acatgaccat ccagnngccg ttctcatt 288

<210> 3161  
<211> 282  
<212> nucleic acid  
<213> Glycine max

<400> 3161

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ttctcagcgc gtaaagcatg gctttgttgg tggagataac cacnctggt ccncgagtac 120  
aagggtcaagg acctttccca ngccgacttc ggccgcctcg agatcgagct ggccgaggtt 180  
gagatgcncg gcctcatggc ctgtcggacc gagttcggcc cctcccagcc cttcaagggg 240  
gcccgcatca ccgntccct ccacatganc atccagaccg cc 282

<210> 3162  
<211> 318  
<212> nucleic acid  
<213> Glycine max  
<400> 3162

cgcacgncgc gaacnnnagc ncgcgaattc ggntcgagng ngccncgac tggggccccg 60  
gtggtggaca ccgacctcat cgtcgacgac ggtggtgang nnacnctnct catccacnaa 120  
ggcgtcaang ccnaggagcn cnatgagaag accggcgaan tcnccgannc caactccacc 180  
ganaacgccg agctgcagat cgnngcttacc atcancagag angggttgaa gaccganccc 240  
atnaggnanc gcaagatgaa ggagcgtctc gttggggtnr ctgaggnnac cancactgga 300  
gttaagaggc tcnatcag 318

<210> 3163  
<211> 319  
<212> nucleic acid  
<213> Glycine max  
<400> 3163

gtcgcangca cgcgtacgta agctcggaat tcggctcgag acggtgcga gaagacgaca 60  
gaaggctcag cttgagttgt ggaaggagaa gagtaccggc aagtacgaga agaaagttta 120  
cgttttgccc aagcaccttg atgagaaggt ggctgcactt caccttgga aacttgagc 180  
taagctcacc aagcttagcc cggcccaggc tgattacatc agtgtgcctg ttgaggggtcc 240  
ataaagcctg ctattacag gtactaagta attgagatta tcaacggaaa gtgagggaaa 300  
gacaaaatcg gttttatga 319

<210> 3164  
<211> 294  
<212> nucleic acid  
<213> Glycine max

<400> 3164

tcgcangcac gcntacgtaa gctcggaatt cggctcgagc ggaaagtgag ggaaagacaa 60  
aatcggtttt atgaatcgga ttgattgttt aattttcctt tgataatctc aattacttag 120  
tacctgtaat gagcaggctt gtatggaccc tcaacaggca cactgatgta atcagcctgg 180  
gccgggctaa gcttgggtgag cttagctcca agtttgccaa ggtgaagtgc agccaccttc 240  
tcatcaaggt gcttgggcaa aacgtaaact ttcttctcgt acttgccggt actc 294

<210> 3165

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 3165

nncgcatgca cgcgtacgta agctcgggaa ttcgggctcg agctctttct ctagtcctgt 60  
tattttctcag cgcgtaaagc atgggctttg ttgggtggaga aaaccacgag tggtcgcgat 120  
acaaggctca ggacctttcc caggccgact tcggccgcct cgagatcgag ctggccgagg 180  
ttgagatgcc cggcctcatg gcctgtncgg accgagttcg gcccctccca gccottcaag 240  
ggggcccgca tcaccggctn cctttccaca tgaccatcca gaccgccgtt ctca 294

<210> 3166

<211> 204

<212> nucleic acid

<213> Glycine max

<400> 3166

cgtcgcangc acgcgtacgt aagctcggaa ttcggtctga ggtttctgag gaaaccacca 60  
ctggagttaa gaggctctat cagatgcagg cgaatgggac tcttctcttc cctgctatta 120  
atgtcaatga ctctgtcacc aagagcaagt ttgacaactt gtatgggtgc cgtcactctc 180  
tccctgatgg tctcatgagg gcta 204

<210> 3167

<211> 203

<212> nucleic acid

<213> Glycine max

<400> 3167

gtcgcangca cgcgtacgta agctcggaat tcggctcgag gtcaccaaga gcaagtttga 60  
 caacttgat ggggtgccgtc actctctccc tgatgggtctc atgaggggcta ccgatgttat 120  
 gattgctgga aaggtggctg ttgtggctgg atatgggtgat gttggcaagg gttgtgctgc 180  
 tgcaatgaag caggttggtg ctc 203

<210> 3168  
 <211> 266  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3168

gcaagtacga gaagaaagtt tacgttttgc ccaagcacct tgatgagaag gtggctgcac 60  
 ttcaccttgg caaacttggg gctaagctca ccaagcttag cccggcccag gctgattaca 120  
 tcagtgtgcc tgttgagggg ccatacaagc ctgctcatta caggtactaa gtaattgaga 180  
 ttatcaacgg aaagtgaggg aaagacaaaa tcggnnttat gaatcggatt gattgtttaa 240  
 ttttcctttt tttgaatttt tgttgt 266

<210> 3169  
 <211> 326  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3169

ntctnntgna ngcgtacgta agctcggaat tnnctcgag ntcgagcngc gccgtcttcn 60  
 anntggacag ggtgagaccc tccagganct actgggtggg caccgngcgc gccctcgact 120  
 ggggccccgg tgggtggaccc gacctcatcg tnnacgacgg tggtgacgct accctttctca 180  
 tccacgaagg cgtcaaggcc gaggagctct ntgagaagac cggcgaattc ccgancccaa 240  
 ntccaccgac aagccggant ttcagatcgt gnttancatc atcagagatg gttgaagacc 300  
 gttccaacca ggttacngca gatgaa 326

<210> 3170  
 <211> 315  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3170

natcgatgca cgcgtacgta agctcggnat tcggctcgan ctcgagccga atcggtcga 60  
 gggttgacca catgaagatn atganganca atgcnattgt anncaacatt ggnncacttt 120  
 natcatnagn tcgacatnct nggggtggag nactaccccc gcgtgangcg catccaccat 180  
 caagccccaa accgacagat ggggtcanccc cgagaccaat gtcggcatca ttgtcttggc 240  
 cgaggggtcgt ttgatgaact tgggatgcgc cacaggacac cctagttttg tgatgtctgt 300  
 cctcacnaac caggt 315

<210> 3171  
 <211> 274  
 <212> nucleic acid  
 <213> Glycine max

<400> 3171

ncgcgtgnac gcgtacgtaa gctcggaatt cggctcgagg tttcctcacc actccctcca 60  
 ttctctttct ctagtcctgt tatttctcag cgcgtaaagc atggctttgt tgtcnggaga 120  
 anaccacgag tggtcgagag tacaaggtca aggaccttc ccaggccgac ttcgcccgcc 180  
 tcgagatcga gctggccgag gttgngatgc ccggcctcat ggcntgtcgg accaggttcg 240  
 gccctccca gcccttcaag ggggcccgc tcaac 274

<210> 3172  
 <211> 282  
 <212> nucleic acid  
 <213> Glycine max

<400> 3172

gtgcgancga cgcgtncgga ngcacgctcn ctttgcctta gtgcctgtta tttctcanen 60  
 cgtaaagcat ggctttgttg gtggagaaaa ccacnagtgg tgcgcgagta canggtinnag 120  
 gacctttgcc caggccgact tcnccgcct cgagatcgag ctggccgagg ttganatgcc 180  
 cggcctcatg gcctgttcgg accgagttcg gccctccca ncccttcaag ggggcccgc 240  
 tcaccggctc cctccacatg accatccaga ncgcggttct ca 282

<210> 3173  
 <211> 312  
 <212> nucleic acid  
 <213> Glycine max

<400> 3173

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cttgntgaga aggtggctgc acttcacctg ggcaaacttg gngctaagct gaccagctt 120  
agcaagtccc aggttgatta catcagtgtg cctgttgagg gtccatacaa gcctgtcac 180  
tacaggctact aagtgattga gatgatcaac tgaaaagtga gtgagggaaa gacaaaaatc 240  
ggttttatca atcggatttg attgtttaat tttccttttt tgatttttgg tgtagactt 300  
tcagatttgn gg 312

<210> 3174

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 3174

angnacangc acgcgtacgt aagctcgga ttcggctcga gtgccaagc accttgatga 60  
gaaggtggct gcaattcacc tgggcaaact tggngctaag ctgaccacgc ttagcaagtc 120  
ncaggctgat tacatcagtg tgctgttga gggccatac aagcctgtc actacaggta 180  
ctaagtgatt gagatgatca actgaaaagt gaggtaggga aagacaaaaa tcggttttat 240  
caatcggatt tgattgttta attttccttt ttttgatttt tgggtgtaga cttttca 297

<210> 3175

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 3175

tcgcntgcac gcgtacgtaa gctcgggaatt cggctcgagt ttacgttttg cccaagcacc 60  
ttgatgagaa ggtggctgca cttcanctgg gcaaacttgg acctaagctg acccagctta 120  
gcaagtccca ggctgattac atcagtgtgc ctgttgaggg tccatacaag cctgtctact 180  
acaggctacta agtgattgan atgntcaact gaaaagtga tgagggaaaag acaaaaaatcg 240  
ntttntcaa tcggatttga ttgtttaatt ttcctttttt tgatttttgg tgtaga 297

<210> 3176

<211> 289

<212> nucleic acid

<213> Glycine max

<400> 3176

gtcgcacgca cgcgtacgta agctcggaat tcggctcgag tgcccaagca ccttgatgag 60  
aagggtggctg cacttcacct gggcncactt ggngctaagc tgacccagct tgcncaaatc 120  
ccaggctgat tacatcagtg tgcctgttga ggggtccatac aagcctgctc actacaggta 180  
ctaagtgatt gagatgatca actgaaaagt gaggtaggga aaggacaaaa atcgggtttta 240  
tcaatcggat ttgatgttta attttccttt tttgattttg gtgttagan 289

<210> 3177

<211> 336

<212> nucleic acid

<213> Glycine max

<400> 3177

gngangcagn gtacgtaagc tcggaattcg gctcgaggag agagagagag agagagagag 60  
atctatctat ctatcaagat ngcggtgttg gttgaaaaaa aaaannattg anaggganta 120  
caagggtgaag ganatgatgc aagccgnttt nggaagattg gaaattcgag ctggcgagg 180  
ttgaaatgcc cggcctcatg tccnccgc accgagttcg gcccctcttc aatccttcaa 240  
gggcgctagg atcancggct cctccacat gaccatcnan agccgncgtc cttcatngag 300  
acnctaccg ctctcggcgc cgaggtcgc tggtgc 336

<210> 3178

<211> 209

<212> nucleic acid

<213> Glycine max

<400> 3178

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ccgagcgcg cctcgactgg ggcncggtg gtggaccga cttcatcgt cgacgacggt 120  
ggtgacgnta cccttctcat ccacgaaggc gtcaaggncg agganctcta tgagaanacc 180  
ggcgaactcg ccganccan ctccacaaa 209

<210> 3179

<211> 291

<212> nucleic acid

<213> Glycine max

<400> 3179

nagtcgcang cacgcgtacg taagctcgga attcggctcg aggctcacca actcccgtc 60  
ccatttcctt atttatagac agagtctgat tgtttcctca ccactccctc cantctcttt 120  
ctcctagtcc tgttatttct cagcgcgtaa agcatggctt tgttggtgga gaaaaccacg 180  
agtggctcgc agtacaaggt caaggacctt tcccaggcgc attcgggcgc cctcgagatc 240  
gagctggcgc aggttgagat gcccggcctc atggcctgtc ggaccgagtt c 291

<210> 3180

<211> 297

<212> nucleic acid

<213> Glycine max

<400> 3180

nacgtcgcat gcacgcgtac gtnagctcgg aattcggctc gagtngaaag accggggaac 60  
tccccgaccc taactccact gacaacgcgc agttccagat cgtgcttacc atcatacaga 120  
gatgggttga agaccgatcc caccaggtac cgcaagatga aggagcgtct cgttgggggtt 180  
tctgaggaaa ccaccactgg tgtaagagg ctatatcaga tgcaggcgat tgggntntat 240  
ttccccgctna taataatnnc nngnnntctg ttaccaagng cngtnnaca acttgnc 297

<210> 3181

<211> 208

<212> nucleic acid

<213> Glycine max

<400> 3181

gtcgcangca cgcgtacgta agctcggaat tcggctcgag tacgttttgc ccaagcacct 60  
tgatgagaag gtggctgcac ttcacctggg caaacttggg gctaagctga cccagcttag 120  
caagtcccag gctgattann ncagtgtgcc tgttgagggt ccatacaagc cgctcactac 180  
aggtactaag tgattgagat gatcaact 208

<210> 3182

<211> 212

<212> nucleic acid

<213> Glycine max



<400> 3182

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctctctttct ctagtcctgt 60  
tattttctcag cgcgtaaagc atggccttgt tggaggagaa aaccacnagt ggtcgcgagt 120  
acaagggtcaa ggacctttcc caggccgact tcggcngcct cgagatcnag ctggccgagg 180  
ttgagatgcc cggcctcatg gctgtcgga cc 212

<210> 3183

<211> 317

<212> nucleic acid

<213> Glycine max

<400> 3183

aagtncncat gcaagctac gtaantcgga attcggctcg agctctagtc ctgttatttc 60  
tcanccgcta aagcatgggc tttgttggtg gagaaaacca cgagtngtcc gctagtacaa 120  
ggtcaaggac ctttcccagg ccgacttcng ccgcctcgag atcgagctgg ccgagggtga 180  
natgcccggc ctcatggcct gtnggaccga ntccggcccc ttcccaaccc ttcaaggggg 240  
cccgatcan cggctccctn canatganca tccagaancg cgttntcatt gngaccctna 300  
ncggctttgg ggcggag 317

<210> 3184

<211> 294

<212> nucleic acid

<213> Glycine max

<400> 3184

tcgcangcag ncgtacgtaa gctcggaatt cggctcgang anaggtgnct gcacttcacc 60  
tgggcaaact tggcnctaag ctgaccanc ttagcaagtc ccaggctgat tacatcagtg 120  
tgcncgttga gggccatac aagcctgctc antannggta ctaagtgatt gagatgatca 180  
actgaaaagt gagtgaggga aagacaaaaa tcggttttat caatcggatt tgattgttta 240  
attttccttt tttgattttt ggtgttngac ttttcagaat gtggtagaag aatt 294

<210> 3185

<211> 245

<212> nucleic acid

<213> Glycine max

<400> 3185

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gcaagccgnt ttcggaagat tggaaatcga gctggcggag gttgaaatgg cccggcatca 120  
tgtccctccc ggcaccngt tcggcccctc ttcaaccctt caagggcgcn angatnaccg 180  
gntccctcca caatgagcnn ncaaanagcc gtacctnaaa cgnagacncg cacnngccng 240  
ggggc 245

<210> 3186

<211> 234

<212> nucleic acid

<213> Glycine max

<400> 3186

aaaannanan gtngcatgca cgcgtacgta agctcggaat tcggctcgnn ctcgagccga 60  
atcggtcga ctttctctag tctgtttatt tctcagcgcg taaagcatgn ctttgttggt 120  
ggagaaaaca nacgagtggg cgcgagtaca aggtcaagga cctttcccag gccgacttcg 180  
gccgntcga gatcgagctg gccgaggttg agatgcccg cctcatggcc tgtn 234

<210> 3187

<211> 298

<212> nucleic acid

<213> Glycine max

<400> 3187

tcgcnngcac gcgtacgtna gctcgnantt cggcnccgag tggnaggagg taaggctggg 60  
tcgaccaga tctagttgag ctcaccaact cccgctccca tttccttatt tatagacaga 120  
gtctgattgt ttctcacca ctccctccan tctctttctc tagtcctgtt atttctcagc 180  
gcgtaaagca tggctttgtt ggtggagaaa accacgagtg gtcgcgagta caaggtcaag 240  
gacctttccc aggccgactt cggccgcctc gagatcgagc tggccgaggt tgagatgc 298

<210> 3188

<211> 221

<212> nucleic acid

<213> Glycine max

<400> 3188

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 cgctcgagcc ggtcctgtna tntctcagcg cgtaaagcat ggctttantt ggtgganaaa 120  
 accacgagtg gtgcgcagta caaggtcaag gacctttccc aggccgactt cgggccccctn 180  
 cgagatcgag ctggccgagg ttgagatgcc cggcctcatg g 221

<210> 3189  
 <211> 291  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3189

anncananaa tnatgcacgc gtacgtaagc tcggaattcg gctcgagatt gtttcctcac 60  
 cactccctcc antctctttc tctagtcctg ttattttctca gcgcgtaaag catggctttg 120  
 ttggtggaga aaaccacgag tggtcganag taanaaggctc aaggactttc ccaggccgac 180  
 ttcgnggcc tcgagatcga gctggccgag gttnaaatgc cgggcctcat ggctggncgg 240  
 acgattnggg cccctcnaa cctttaaggg gggccnaaat cangggntcc n 291

<210> 3190  
 <211> 303  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3190

ncgcgatgca gcgtacgtaa gctcggaatt cggctcgagg ttattttctca gcgcgtaaag 60  
 catggctttg tnggtggaga aaaccacgag tggtcgcgag tacaagggtca aggacctttc 120  
 ccaggccgac ttcgcccgcc tcgagattcg agctggncgg aggttgagat gcccgacct 180  
 catggcctgt ncggaccgag ttnggncccc taccagccc tttcaagggg gncccgcatc 240  
 accggcnccc nccacatgna ccatccagt cgnccgttg ttcattgn gn accctgcacc 300  
 gcc 303

<210> 3191  
 <211> 144  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3191

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ngcaggcacg cgtacgtaag ctncggaatt cggctcgagn cggctcgagg ggttggtgctg 60
ctgcattgaa gcaggctggg gctcgtgtca tegtgactga gattgacccc atttggtgccc 120
ttcaggctct catggaaggg cctt 144

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<210>      3192
<211>      134
<212>      nucleic acid
<213>      Glycine max

<400>      3192

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cgacggtggg gacgctaccc ttctcatgcc acgaaggcgt ttaggccgag gagctctatg 120
agaagaccgg cgaa 134

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<210>      3193
<211>      303
<212>      nucleic acid
<213>      Glycine max

<400>      3193

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ctgattacat cagtgtgcct gttgaggggc catacaagcc tgctcactac aggtactaag 120
tgattganat gatcaactga aaagtgagtg agggaaagac aaaaatcggg tttatcaatc 180
ggatttgatt gtttaatttt cctttttttg atttttgggt ttagactttt cagatttgtg 240
gtagaagaat gtagccattt ttatttctgt agaacttttg ttcggtgggt gggaccagta 300
agg 303

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<210>      3194
<211>      315
<212>      nucleic acid
<213>      Glycine max

<400>      3194

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gtgtgccacg ggacacccca gctttgtgat gtcgtgctcc ttcaccaacc aggtcatngc 120
tcagcttgaa ttgtggaaag agaagggttc tgggaagtat gagaagaagg tgnatgtgtt 180

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gccaagcac cttgacngga aagtggcttc tctccacctt ggccagnttg gagctaggct 240  
 caccaagctt tccanagacc aagctgatta catcagtgtg cctgttgagg gtccatacaa 300  
 gccgctccnc acagt 315

<210> 3195  
 <211> 290  
 <212> nucleic acid  
 <213> Glycine max

<400> 3195

cancacactc gcangcacgc gtacgtaagc tcggaattcg gtcgagcag gtcattgctc 60  
 agcttgaatt gtggaaagag aagggttctg ggaagtatga gaagaagggtg tatgtgttgc 120  
 ccaagcacct tgacgagaaa gtggcttctc tccaccttgg ccagcttgga gctaggctca 180  
 ccaagctttc caaagaccaa gctgattaca tcagtgtgcc tgttgagggt ccatacaagc 240  
 ctgctcacta caggtactga tccatcctat tgggggagaa taaacctaaa 290

<210> 3196  
 <211> 217  
 <212> nucleic acid  
 <213> Glycine max

<400> 3196

gtcgcangca cgcgtacgta agctcggaat tcggctcgag ctcagcttga attgtggaaa 60  
 gagaagggtt ctgggaagta tgagaagaag gtgtatgtgt tgccaagca ccttgacgag 120  
 aaagtggctt ctctccacct tggccagctt ggagctaggc tcaccaagct ttccaaagac 180  
 caagctgatt acatcagtgt gcctgttgag ggtccat 217

<210> 3197  
 <211> 255  
 <212> nucleic acid  
 <213> Glycine max

<400> 3197

gaaagagaag ggttctggga agtatgagaa gaagggtgat gtgttgccca agcaccttga 60  
 cgagaaagtg gcttctctcc accttgcca gcttgagct aggtcacca agctttccaa 120  
 agaccagctg attacatcag tgtgctgttg angggggcca taanagcttg tcnctnangg 180

nnnnngnccn ncctttgggg gggaannaac ccgaantntn ttnattcgg ggggggnttg 240  
tnnantttnn ttng 255

<210> 3198  
<211> 338  
<212> nucleic acid  
<213> Glycine max  
<400> 3198

aggaccatgc cgccgcgcgc atcgcccgcg acaggcctcc gtcttcgcct ggaagggatga 60  
gaccctccag gaatactggt ggtgcaccga gcgcgcctcg actgggnncc ccggcgccgg 120  
cccgatctca tcgtcgacga cggcgggcgac accactcttc tcattcacga gggcgtcaag 180  
gccgaggaga tctttgagaa gaccggccag tccccgacc cggcttcctc cgacaatgcg 240  
gattncgat cgtgctgagc atcatcaggg atggttgaag accgatccca agaggtacca 300  
caagatgaag acagaatcgt cgggtgtctcc gaagaaac 338

<210> 3199  
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<212> nucleic acid  
<213> Glycine max  
<400> 3199

agtgcacatgc acgcgtacgt aagctcgga ttcggctcga ggcttgaag ggtgagaccc 60  
tccaggaata ctggtggtgc accgagcncg cctcgactg gngccccggc ggcgcccccg 120  
antcatcgt ccgacgacgg cggcgacacc actcttctca ttcacgaggg cgtcaaggcc 180  
gaggagatct ttgagaagac cggccagttc cccgaccggg ctctctccga caatgcggag 240  
ttccagatcg tgctgagcat cattcagggg tggcttgaag accgatccca agaggtacca 300  
caagatgaag gacagaa 317

<210> 3200  
<211> 290  
<212> nucleic acid  
<213> Glycine max  
<400> 3200

gtcgccangca cgcgtacgta agctcggaat tcggctcgag ggaatatcca ctagcttcgt 60

ggaggtgacg gatcttgaca tggttgatgc tncatagta gaagggaaaa caaaagtggc 120  
 tttacttcga atctgtttcc aacccccacc ttacgggttc gaacatacct gaactgtgcc 180  
 acatggcaca ccggaaggga gtgacggtgg tggtaggaca cacgttcgcg cccatggtgc 240  
 tttcgccagc gcgtcttggt gctgatgttg tegtacacag tatctccaag 290

<210> 3201  
 <211> 213  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3201

attcggctcg aggcggaata tccactagct tcgtggaggt gacggatctt gacatggttg 60  
 atgctgccat agtagaaggg aaaacaaaag tgctttactt cgaatctgtt tccaacccca 120  
 cccttacggt tgcgaacata cctgaactgt gccacatggc acaccggaag ggagtgacgg 180  
 tggtaggtgga caacacgttc gcgcccatgg tgc 213

<210> 3202  
 <211> 297  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3202

cncangcacg cgtacgtaag ctcggaattc ggctcgaggt gggacccacg cgctcctctc 60  
 acactttctc ccgcgcacgt gcggaatatc cactagcttc gtggaggtga cggatcttga 120  
 catggttgat gctgccatag tagaaggga aacaaaagt ctttacttgc aatctgtttc 180  
 caacccacc cttacggttg cgaacatacc tgaactgtgc cacatggcac accggaaggg 240  
 agtgacggtg gtggtggaca acacgttcgc gcccatggtg ctttcgccag cgcgtct 297

<210> 3203  
 <211> 300  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3203

gtcncnngta cgtaanctcg gaattcggct cgagcngaca ancccaann ccaagcccaa 60  
 caatctgcat ccccgccgc ggccgtgca accaaatggg ccgtggacag ctggaagtcc 120

aagaaggccc tgcagntgcc cgaatacccc aaccaggagg atctcgaggc cgtcctccgc 180  
 accctcgacg cntnccccctc anatcgtctt cgcggcgag gcccgganac tcgaggagca 240  
 cctcgccgag gccgccatng gaaatgcntt cttecnan ggcgnagatg tnccnagagt 300

<210> 3204  
 <211> 434  
 <212> nucleic acid  
 <213> Glycine max  
 <400> 3204

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ccacgcgtcc gccacgcgt cgggtccgcc atctccgcc tectnctcca gctctgcagc 60  
 cacngcgaac anggtggtcg nctccagaac tctntacngt gggacccacn cgtcctctc 120  
 acactttctc ccgcgcacgt gcggaatata cactagcttc gtggaggtga cngatcttga 180  
 catggnttat gctgccataa tagaaaggaa aacaaaagtg ctttacttnt aatctggttc 240  
 caaccccacc cttacngttg cgaacatacc tgaactgtgc cacatggcac accggaaggg 300  
 agtgactgtg gtggtggaca acacgttcgt gcccatgggtg ctttcgccag cgcgtntttg 360  
 gtgcttatgt ttgtnccttca cagtatctcc aagttcatna atnggtgggg cccgatatta 420  
 ttgcangagc ggng 434